

APPLICATION
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TITLE: METHODS AND COMPOUNDS FOR PREVENTION OF
GRAFT REJECTION

APPLICANT: TERRY B. STROM AND TOWIA LIBERMANN

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METHODS AND COMPOUNDS FOR PREVENTION OF GRAFT REJECTION

This application is a continuation-in-part of pending U.S. patent application Serial No. 08/024,569, filed
5 March 1, 1993 by Terry Strom, Vicki E. Rubin-Kelley, Towia Libermann entitled "Methods and Compounds for Prevention of Graft Rejection"; which is a continuation-in-part of U.S. patent application Serial No. 07/843,731, filed February 28, 1992, now abandoned.

10 Statement as to Federally Sponsored Research

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15 Field of the Invention

This invention relates to methods and compounds useful in inhibiting the immune response in a mammal.

Background of the Invention

For many life-threatening diseases, organ
20 transplantation is considered a standard treatment and, in many cases, the only alternative to certain death. The immune response to foreign cell surface antigens on the graft, encoded by the major histocompatibility complex (MHC) and present on all cells, generally precludes successful
25 transplantation of tissues and organs unless the transplant tissues come from a compatible donor or the normal immune response is suppressed. The best compatibility and thus, long term rates of engraftment, are achieved using MHC identical sibling donors or MHC identical unrelated cadaver
30 donors (Strom, 1989, *supra*; Strom, 1990, Clinical Aspects of Autoimmunity 4:8-19).

The host response to an organ allograft involves a complex series of cellular interactions among T and B lymphocytes as well as macrophages or dendritic cells that recognize and are activated by a foreign antigen (Strom, 1989, *supra*; Strom, 1990, Clinical Aspects of Autoimmunity 4:8-19). Co-stimulatory factors, primarily cytokines, and specific cell-cell interactions, provided by activated accessory cells such as macrophages or dendritic cells are essential for T cell proliferation. These macrophages and dendritic cells either directly adhere to T cells through specific adhesion proteins or secrete cytokines that stimulate T cells, such as IL-1 and IL-6 (Strom, 1989, *In: Organ Transplantation: Current Clinical and Immunological Concepts*: 4:8-19; Strom, 1990, Clinical Aspects of Autoimmunity 4:8-19).

IL-1 induces expression of the IL-6 gene in accessory cells. Accessory cell-derived co-stimulatory signals stimulate activation of interleukin-2 (IL-2) gene transcription and expression of high affinity IL-2 receptors in T cells (Pankewycz, et al., 1989 Transplantation 47:318; Cantrell, et al., Science 224:1312; Williams, et al., 1984 J. Immunol. 132:2330-2337). IL-2, a 15 kD protein, is secreted by T lymphocytes upon antigen stimulation and is required for normal immune responsiveness. IL-2 stimulates lymphoid cells to proliferate and differentiate by binding to IL-2 specific cell surface receptors (IL-2R). IL-2 also initiates helper cell activation of cytotoxic T cells and stimulates secretion of γ -interferon (γ -IFN) which in turn activates cytodestructive properties of macrophages (Farrar, et al., 1981 J. Immunol. 126:1120-1125). Furthermore, γ -IFN and IL-4 are also important activators of MHC class II expression in the transplanted organ, thereby further expanding the rejection cascade by actually making the

grafted organ more immunogenic (Pober, et al., 1983, J. Exp. Med., 157:1339; Kelley, et al., 1984 J. Immunol., 132:240-245).

5 Similar mechanisms are involved in the development
of autoimmune disease, such as type I diabetes. In humans
and non-obese diabetic mice (NOD), insulin-dependent
diabetes mellitus (IDDM) results from a spontaneous T-cell
dependent autoimmune destruction of insulin-producing
10 pancreatic β cells that intensifies with age. The process
is preceded by infiltration of the islets with mononuclear
cells (insulitis), primarily composed of T lymphocytes
(Bottazzo, G.F., et al., 1985, J. Engl. J. Med., 113:353;
Miyazaki, A., et al., 1985, Clin. Exp. Immunol., 60:622). A
delicate balance between autoaggressive T-cells and
15 suppressor-type immune phenomena determine whether
expression of autoimmunity is limited to insulitis or
progresses to IDDM. In NOD mice, a model of human IDDM,
therapeutic strategies that target T-cells have been
successful in preventing IDDM (Makino, et al., 1980, Exp.
20 Anim., 29:1). These include neonatal thymectomy,
administration of cyclosporine A, and infusion of anti-pan
T-cell, anti-CD4, or anti-CD25 (IL-2R) mAbs (Tarui, et al.,
1986, Insulitis and Type I Diabetes. Lessons from the NOD
Mouse. Academic Press, Tokyo, p. 143).

25 The aim of all rejection prevention and reversal
strategies is to suppress the patient's immune reactivity to
the graft, with a minimum of morbidity and mortality.
Existing immunosuppressive therapies include administration
of immunosuppressive compounds such as cyclosporine A, FK506
30 and rapamycin (First, 1992 Transplantation, 53:1-11).
Because these agents inhibit proliferation of T cells
generally, systemic treatment of patients leads to systemic
immunosuppression which carries with it potential

complications, including increase risk of infections and cancer (Wilkinson, et al., Transplant, 47:293-296; Penn, 1991 Transplant Proc., 23:1101; Beveridge, et al., 1984 Lancet, 1:788). In addition, these immunosuppressive agents

5 cause considerable side effects, including nephrotoxicity, hepatotoxicity, hypertension, hirsutism, and neurotoxicity (Strom, 1989, *supra*; Strom, 1990, Clinical Aspects of Autoimmunity, 4:8-19; Tilney, et al., 1991 Ann Surg., 214:42-49; Myers, et al., 1984 N. Engl. J. Med., 311:699).

10 Administration of monoclonal antibodies against the T cell-specific antigen CD3 has been shown to block acute allograft rejection (Mackie, et al., 1990 Transplantation, 49:1150). Antibodies directed against the IL-2 receptor, T cell receptor, CD4, and certain cell adhesion molecules such

15 as ICAM-1 have been used (Cosimi, et al., 1990 Surgery, 108:406; Cosimi, et al., 1990 J. Immunol., 144:4604; Strom et al., 1989 Kidney Int., 35:1026; Walz, 1990 Transplantation, 49:198-201). Anti-IL-2 receptor antibodies have been shown to bring about improved patient and graft

20 survival (Soulillou, et al., 1990 N. Engl. J. Med., 322:1175).

Two novel cytokines, TGF- β and IL-10, have recently been identified. These proteins have immunosuppressive activity, and apparently act via different mechanisms on the

25 immune system (Moore, et al., 1990 Science, 248:1230-1234; Vieira, et al., 1991 Proc. Natl. Acad. Sci USA, 88:1172-1176; Barnard, et al., 1990 Biochim Biophys. Acta, 1032:79-87; Derynck, et al., 1985 Nature, 316:701-705).

TGF- β exhibits potent immunosuppressive effects

30 including inhibition of B and T cell activation and differentiation and deactivation of activated macrophages (Barnard, et al., 1990 Biochim Biophys. Acta, 1032:79-87; Roberts et al., 1990 Handbook of Experimental Pharmacology,

95:419-472). In particular, TGF- β inhibits expression of immunoglobulin genes in B cells, decreases IL-2 induced T cell proliferation and differentiation of cytotoxic T cells, and inhibits MHC class II antigen expression on a number of cell types (Kehrl, et al., 1986 J. Exp. Med., 163:1037-1050; Ruegamer, et al., 1990 J. Immunol., 144:1767-1776; Cross, et al., 1990 J. Immunol., 144:432-439; Nelson, et al., 1991 J. Immunol., 146:1849-1857; Lee, et al., 1987 J. Exp. Med., 166:1290-1299; Wright, et al., 1986 Diabetes, 353:1174-1177). TGF- β has been shown to inhibit pancreatic islet allograft rejection in mice, suggesting its potential use as an immunosuppressive agent. TGF- β is produced by many different cell types, including macrophages, B and T cells, lung and mesenchymal cells, skin cells, platelets, and bone cells (Barnard, et al., 1990 Biochim Biophys. Acta, 1032:79-87; Roberts et al., 1990 Handbook of Experimental Pharmacology, 95:419-472). In addition, some cancer patients have been observed to show signs of immunosuppression due to secretion of TGF- β by the cancer cells (Barnard, et al., 1990 Biochim Biophys. Acta, 1032:79-87; Roberts et al., 1990 Handbook of Experimental Pharmacology, 95:419-472; Siepl, et al., 1988 Eur. J. Immunol., 18:593-600).

IL-10 was first identified as a product of activated T_H2 T helper cells with the ability to inhibit macrophage-dependent cytokine synthesis in T_H1 T helper cells (Fiorentino, et al., 1989 J. Exp. Med., 170:2081). IL-10 appears to be expressed by several different hematopoietic cell types, including activated T_H2 cells, activated macrophages, mast cells, and B cells (Moore, et al., 1990 Science, 248:1230-1234; O'Barra, et al., 1990 Int. Immunol., 2:821; De Waal Malefyt, et al., 1991 J. Exp. Med., 174:1209-

1220). IL-10 appears to inhibit the expression of a number of cytokines in macrophages, including interleukin-6 (IL-6), interleukin-1 (IL-1), interleukin-8 (IL-8), tumor necrosis factor- α (TNF- α), granulocyte/macrophage colony stimulatory factor (GM-CSF) and granulocyte colony stimulatory factor (G-CSF). IL-10 diminishes the antigen-presenting capacity of macrophages via downregulation of MHC class II gene expression on macrophages, and induces expression of MHC class II genes, but not class I genes in unstimulated splenic B cells (Go, et al., 1990 J. Exp. Med., 172:1625-1631; De Waal Malefyt, et al., 1991 J. Exp. Med., 174:915-924).

Summary of the Invention

We have developed novel methods for achieving localized immunosuppression. These methods may be used for the prevention of graft rejection following organ or tissue transplantation and for the prevention of tissue destruction due to autoimmune disease. Our methods allow strong local immunosuppression without the side effects associated with general immunosuppressive methods now in use. In addition to these methods, we have discovered a new immunosuppressive protein.

Accordingly, in one aspect, the invention features a method for inhibiting rejection of a transplanted tissue in a mammal, involving (a) introducing into a cell, either in vivo or ex vivo, DNA encoding an immunosuppressive protein such that the cell expresses (and preferably secretes) the immunosuppressive protein. If the DNA is introduced ex vivo, the method further includes transplanting the cell into the mammal. The cell may be a cell of the transplanted tissue or organ or may be a cell to be transplanted to the region of the body near the transplanted tissue organ. The immunosuppressive protein is, according to the invention,

produced in the localized anatomical region where it is required, i.e., in the vicinity of the transplanted tissue, but is not administered systemically to the animal. Thus, generalized systemic immunosuppressive effects are not produced. This localized expression is achieved by providing the DNA encoding the immunosuppressive protein under the regulatory control of a promoter sequences not naturally found immediately 5' to the immunosuppressive polypeptide-encoding DNA. The promoter may be constitutive, tissue-specific, immune response inducible, or inducible by orally or systemically administered compounds.

The method can be used to inhibit rejection of both allografts and xenografts, e.g., transplanted organs such as heart, kidney, liver and lung, and tissues such as bone and skin, or cellular transplants, e.g., islets.

In a second aspect, the invention features a method for inhibiting tissue damage due to autoimmune disease in a mammal by introducing into a cell, either in vivo or ex vivo DNA encoding the immunosuppressive protein such that the cell expresses (and preferably secretes) the immunosuppressive protein. Where the DNA is introduced ex vivo, the method further includes transplantation of the tissue into the mammal. Using either method of introduction, the immunosuppressive protein is expressed (and preferably secreted) by the cell of the tissue at levels sufficient to produce an immunosuppressive effect. The local expression of the immunosuppressive protein is achieved by providing the DNA encoding the immunosuppressive protein under the regulatory control of a promoter not naturally found immediately 5' to the DNA encoding the immunosuppressive protein. The immunosuppressive protein is, according to the invention, produced in the localized anatomical region where it is required, i.e., in the

vicinity of the diseased or disease prone tissue or organ, but is not administered systemically to the animal, so that generalized systemic immunosuppressive effects are not produced.

5 The method can be used to decrease autoimmune damage to organs such as heart, kidney, liver, and lung, and to tissues such as bone, skin, neuronal tissue, and synovium.

Both the methods of transplant rejection inhibition and the inhibition of autoimmune related tissue destruction
10 involve localized suppression of the immune response. These methods can employ DNA which encodes any immunosuppressive protein. Examples of suitable proteins are interleukin-10 (IL-10), transforming growth factor β (TGF- β), and the suppressor factor described below produced by human T-cell
15 clones such as IS-2.15. Also included are genes encoding immunosuppressive fusion proteins, such as IL-2Ig, CD2Ig, CTLA-4Ig wherein the N-terminus includes amino acids present in human IL-2, CD-2 or CTLA-4 and the C-terminus includes amino acids present in the Fc fragment of the Ig heavy
20 chains. These proteins all directly decrease the immunosuppressive response in the region where they are expressed.

In a third aspect, the methods of the invention can also employ DNA encoding a protein which indirectly acts as
25 an immunosuppressant. For example, the method may employ the use of a transgene which encodes cyclosporine synthetase. Thus, the cyclosporine synthetase locally processes cyclosporine precursor molecules into the immunosuppressant compound, cyclosporine. In another
30 embodiment, DNA encoding a polypeptide which metabolizes cyclophosphamide to the active form may also be provided using the methods of the invention. Where DNA encoding a protein which processes a prodrugs to yield active

immunosuppressant is employed, the method also features the subsequent administration of the prodrug. As with directly immunosuppressive polypeptides, use of a DNA construct expressing a peptide which acts indirectly creates local immunosuppression without the systemic side effects of general immunosuppression.

In a fourth aspect, DNA encoding a glycosidase may be provided to cells in the region where immunosuppression is desired. Synthesis of such a protein will locally remove antigenic determinants which are targets for xenoreactive natural antibodies.

Expression of directly or indirectly immunosuppressive proteins or glycosidases in the cells in or near the transplanted tissue or in the area of autoimmune response can be controlled by regulatory sequences which cause constitutive expression, or alternatively, expression can be controlled by regulatory sequences which are inducible; in one preferred embodiment, the regulatory sequence controlling expression of the immunosuppressive protein is inducible by a compound which stimulates an immune response. For example, the promoter can be inducible by IL-1, which is expressed during activation of responses. Thus, responses that cause rejection of the transplanted organ or destruction of the autoimmune target tissue when the immune system is being stimulated to cause a rejection turns on the gene for the immunosuppressive protein and halts or slows rejection. Alternatively, the promoter controlling transcription of the immunosuppressive protein gene is inducible by a foreign antigen of the transplanted tissue itself; in this case, as well, the same component which tends to cause rejection also turns down expression of the protein inhibiting rejection.

Promoters which may be used for the expression of the immunosuppressive proteins or glycosidases of the invention are shown in Table 1. In one embodiment, the regulatory sequence controlling expression of the immunosuppressive protein contains a binding site for a non-mammalian (e.g. bacterial) inducible regulator. In this embodiment DNA encoding constructively expressed non-mammalian regulatory protein is also provided. The DNA encoding the regulatory protein may be provided as a part of the DNA sequences encoding the immunosuppressive protein or they may be provided in a second construct. In this embodiment, the inducer is then provided separately, preferably orally, as needed to prevent graft rejection or autoimmune mediated tissue destruction. Thus, the positive bacterial regulatory gene NAHR encodes a factor which activates the PG promotor only in the presence of Na-Salicylate (Yen. J. Bacteriol., 173:5328-35 (1991), incorporated herein by reference).

Preferred promotor-gene combinations for the treatment of a graft rejection of various tissues are shown in Table 2, top. Preferred promoter-gene constructs for the treatment of specific autoimmune diseases are shown in Table 2, bottom. In addition, constructs are shown in Figs. 10-15. These constructs may be used to express immunosuppressive proteins other than those specifically depicted.

The methods of the invention may be used alone for the treatment of autoimmune disease or graft rejection or they may be used in combination with other immunosuppressive therapies.

Constructs of the invention which are immunosuppressive, as used herein, are those constructs which prolong engraftment beyond the 50% rejection period

for the control animal similarly treated, but lacking the transgene. For example, constructs which include non-specific constitutative, immune-responsive, inducible, or pancreatic β -cell specific promoters may be tested in the pancreatic islet cell transplant model described below.

Tissue specific expression may be tested in transgenic mice containing the appropriate promotor constructs. Once the mouse is constructed, expression is tested by reverse transcriptase polymerase chain reaction and Northern blot hybridization using RNA and DNA isolated from different mouse tissues and organs. Constructs for use in the treatment of autoimmune disease are also defined as those which show at least a 20% decrease in tissue damage relative to a control lacking the therapy over a period of time required for complete disease progression in the appropriate mouse model. For example, since with or without the *lpr* gene, NOD mice and EAE model mice may be used to choose among constructs for use in the methods of invention. These mice are already publicly available from suppliers such as Jackson Labs, Bar Harbor, Maine. Useful constructs are also those which provide a mixed lymphocyte reaction (MLR) by decreasing proliferation by 20%, more preferably 40%, and most preferably, by 60% relative to control cells not expressing the transgene. This assay may be done as described in *Current Protocols in Immunology* (1992, Wiley & Sons, Eds., Coligan et al.).

In a fifth aspect, the invention features a substantially pure suppressor factor protein or biologically active analog or fragment thereof. The protein is characterized in that it is secreted by cloned anergic T-cells (e.g., IS-2.15 cells), it blocks interleukin 2 (IL-2)-stimulated T-cell proliferation, it has an apparent molecular weight of between 10 and 30 kilodaltons, it can be

inactivated by heating to 65°C for 15 minutes, it blocks interleukin 4 (IL-4)-stimulated T-cell proliferation *in vitro*, it is non-cytotoxic to T-cells, and it does not inhibit the production of IL-2 by T-cells *in vitro*.

5 "Anergic T-cells", as used herein, refers to T-cells that are hyporesponsive or unresponsive to an antigen or mitogen. By "apparent molecular weight" is meant molecular weight as determined by filtration through microconcentrator tubes with multiple membrane size cut-offs, followed by
10 testing of the resulting concentrates and filtrates for T-cell suppressor activity as described in the detailed description to follow.

The invention also features a purified nucleic acid encoding the new suppressor factor of the invention.

15 The nucleic acid of the invention can be used to alter the effect of IL-2 on cells in a mammal that express, constitutively or transiently, the IL-2 receptor, in order to inhibit IL-2 induced cell proliferation, the proliferation of autoreactive T-cells. The method involves
20 bringing into close proximity of the cell a second cell which is transfected with the nucleic acid encoding the suppressor factor, so that the second cell secretes a protein which causes an alteration of the IL-2 effect. The mammal is preferably a human, and the cell is preferably a
25 T-cell, an endothelial cell lining a blood vessel, or an epithelial cell, most preferably an epithelial cell of the kidney proximal tubule, or an epithelial cell of the gut.

The nucleic acid of the invention can also be used to alter the effect of IL-2 on cells in a mammal that
30 express, constitutively or transiently, the IL-2 receptor. The method involves transfecting the cell with the nucleic acid so that the cell secretes the suppressor factor, causing alteration of the IL-2 effect.

In a related aspect, the invention features a method of altering the effect of IL-4 on cells in a mammal that express, constitutively or transiently, the IL-4 receptor, in order to inhibit IL-4 induced cell proliferation. The method involves bringing into close proximity of the cell a second cell which is transfected with the nucleic acid encoding the suppressor factor, so that the second cell secretes the factor causing an alteration of the IL-4 effect. The mammal is preferably a human, and the cell is preferably a T-cell, an endothelial cell lining a blood vessel, or an epithelial cell, most preferably an epithelial cell of the kidney proximal tubule, or an epithelial cell of the gut.

In another aspect, the nucleic acid of the invention can be used in a method of altering the effect of IL-4 on cells in a mammal that express, constitutively or transiently, the IL-4 receptor. The method involves transfecting the cell with the nucleic acid encoding the suppressor factor so that the cell secretes the factor causing alteration of the IL-4 effect.

The suppressor factor protein of the invention can be obtained from any suitable naturally occurring source and can also be made recombinantly. Also included in the invention are biologically active fragments and analogs of the suppressor factor. The term "fragment", as applied to a polypeptide, will ordinarily be at least about 10 contiguous amino acids, more typically at least about 20 contiguous amino acids, usually at least about 30 contiguous amino acids, preferably at least about 40 contiguous amino acids, more preferably at least about 50 contiguous amino acids, and most preferably at least about 60 to 80 or more contiguous amino acids in length. Biologically active

fragments of the suppressor factor can be generated by methods known to those skilled in the art.

5 A suppressor factor polypeptide, fragment, or analog is biologically active if it exhibits a biological activity of naturally occurring IS-2.15 suppressor factor, e.g., the ability to block IL-2 stimulated T-cell proliferation. The ability of a candidate analog or fragment to block IL-2 stimulated T-cell proliferation can be assessed by methods known to those skilled in the art, e.g., by methods 10 described below or above, such as the islet transplant rejection assay or the MCR assay.

The invention also includes biologically active analogs of the suppressor factor protein of the invention. 15 Analogs can differ from naturally occurring IS-2.15 suppressor factor by amino acid sequence differences or by modifications that do not affect sequence, or by both. Modifications include *in vivo* or *in vitro* chemical derivatization of polypeptides, e.g., acetylation, or carboxylation. Also included are modifications of 20 glycosylation, e.g., those made by modifying the glycosylation patterns of a polypeptide during its synthesis and processing or in further processing steps, e.g., by exposing the polypeptide to enzymes that affect glycosylation derived from cells that normally provide such 25 processing, e.g., mammalian glycosylation enzymes. Also embraced are versions of the same primary amino acid sequence that have phosphorylated amino acid residues, e.g., phosphotyrosine, phosphoserine, or phosphothreonine.

30 Analogs can differ from naturally occurring IS-2.15 suppressor factor by alterations of their primary sequence. These include genetic variants, both natural and induced. Induced mutants may be derived by various techniques, including random mutagenesis of the encoding nucleic acids

using irradiation or exposure to ethanemethylsulfate (EMS), or may incorporate changes produced by site-specific mutagenesis or other techniques of molecular biology. See, Sambrook, Fritsch and Maniatis (1989), Molecular Cloning: A Laboratory Manual (2d ed.), CSH Press, hereby incorporated by reference. Also included are analogs that include residues other than naturally occurring L-amino acids, e.g., D-amino acids or non-naturally occurring or synthetic amino acids, e.g., β or γ amino acids.

Analog of the invention, to be biologically active, will generally exhibit at least 70%, more preferably 80%, more preferably 90%, and most preferably 95% or 99%, homology with all or part of a naturally occurring IS-2.15 suppressor factor amino acid sequence. The length of comparison sequences will generally be at least about 8 amino acid residues, usually at least 20 amino acid residues, more usually at least 24 amino acid residues, typically at least 28 amino acid residues, and preferably more than 35 amino acid residues.

"Homologous", or "homology", as used herein, refers to the subunit sequence similarity between two polymeric molecules, e.g., between two nucleic acid molecules, e.g., two DNA molecules, or two polypeptide molecules. When a subunit position in both of the two molecules is occupied by the same monomeric subunit then they are homologous at that position. The homology between two sequences is a direct function of the number of matching or homologous positions, e.g., if half, e.g., 5 of 10, of the positions in two compound sequences are homologous then the two sequences are 50% homologous, if 90% of the positions, e.g., 9 of 10, are matched or homologous the two sequences share 90% homology. By way of example, the DNA sequences 3'ATTGCC'5 and 3'TATGGC'5 are 50% homologous.

The invention also includes proteins encoded by DNA that hybridizes to IS-2.15 suppressor factor-encoding nucleic acids, and polypeptides or proteins specifically bound by antisera to IS-2.15 suppressor factor, e.g.,
5 antisera to the active site or binding domain of IS-2.15 suppressor factor.

The invention also includes polypeptides that differ from the naturally-occurring suppressor factor by substitution of one amino acid for another of the same
10 class, or that differ by one or more non-conservative amino acid substitutions, deletions, or insertions, located at positions of the amino acid sequence that do not destroy the biological activity of the polypeptide. Amino acids of the same class are ones that share characteristics of
15 hydrophobicity, charge, pKa, or other conformational or chemical properties (e.g., valine for glycine, arginine for lysine, etc.)

As used herein, the term "substantially pure" describes a protein, e.g., an IS-2.15 suppressor factor
20 protein or polypeptide, that has been separated from components, i.e., the components of a eukaryotic cell. Typically, a protein is substantially pure when at least 60%, more preferably at least 75%, more preferably at least 90%, and most preferably at least 99%, of the total material
25 (by volume, by wet or dry weight, or by mole per cent or mole fraction) in a sample is the compound of interest. Purity can be measured by any appropriate method, e.g., in the case of polypeptides by column chromatography, polyacrylamide gel electrophoresis, or HPLC analysis.

30 A "purified nucleic acid", as used herein, refers to a nucleic acid sequence or fragment that is not associated with the sequences that flank it in a naturally occurring state, e.g., a DNA fragment that has been removed from the

sequences that are adjacent to the fragment, e.g., the sequences adjacent to the fragment in its normal site in the genome. The term also applies to nucleic acids that have been substantially purified from other components that naturally accompany it in the cell, and also applies to cDNA and synthetic nucleic acids.

The properties of the suppressor factor of the invention render it and its biologically active analogs and fragments useful in a number of therapeutic and diagnostic applications. In therapy in particular, the factor, because of its immunosuppressive effects, may be useful in the treatment of autoimmune diseases such as lupus, type 1 diabetes, and rheumatoid arthritis. The factor may also be used to inhibit transplant rejection and graft versus host disease (GVHD) following transplantation. When used for this purpose, the suppressor factor will be administered generally by the same regimens, and in the same dosage range, as current commercially-available immunosuppressive agents, with the provision that, because the suppressor factor of the invention is a protein, it will preferably be administered intravenously rather than orally. A nucleic acid encoding the suppressor factor of the invention may be used to protect cells against autoreactive T-cells.

Other features and advantages of the invention will be apparent from the following detailed description thereof, and from the claims.

Detailed Description

The drawings are first described.

Drawings

Fig. 1. is a Southern blot of products of a reverse transcriptase-polymerase chain reaction (RVT-PCR). RVT-PCR transcripts of IL-2 in islet cell grafts and spleens on day 8 post engraftment. Four mice were treated with DAB486-IL-2

(lanes 1-4) and four with anti-CD3 mAb (lanes 5-8) for 8 days. All harvested tissue was snap frozen in liquid nitrogen and total RNA extracted by the GCN method. Five μ g of total RNA was used as starting material for RVT-PCR.

5 Products were size separated on a 1.5% agarose gel to confirm the product size and blotted to a nylon membrane and probed with a cDNA labelled with 32 P. The blots were then scanned using the PHOSPHOR IMAGER system (Molecular Dynamics Inc.) (NK Normal kidney).

10 Fig. 2 is a Southern blot of PCR products. PVT-PCR transcripts of IFN Gamma in 5 rejecting grafts (lanes 9-13). 4 grafts treated with DAB-IL-2 (lanes 1-4) for 8 days and 4 grafts treated with anti-CD3 mAb (lanes 5-8) for 8 days. All grafts were harvested on day 8 post transplant, snap
15 frozen in liquid nitrogen and processed for total RNA. Two μ g of total RNA was used for RVT-PCR. Products were run on a 1.5% agarose gel and blotted to a nylon filter then hybridized with an IFN Gamma cDNA probe labelled with 32 P. The blots were then scanned using the PHOSPHOR IMAGER system
20 (Molecular Dynamics Inc.) (NK normal kidney. BL blank.)

Fig. 3 is a Southern blot of PCR products. RVT-PCR products of coamplified GRANZYME B and TCR C α mRNA transcripts from spleens 1, 4, 12, 21 days post transplant, in rejecting model of pancreatic islet cell transplantation,
25 and CON A stimulated spleen cells. One μ g of total RNA per sample was used as starting material. The products were size separated on a 2.5% agarose gel, blotted to a nylon membrane and cohybridized with 32 P labelled cDNA probes for GRANZYME B and TCR C α . Using the PHOSPHOR IMAGER and
30 IMAGEQUANT software (Molecular Dynamics Inc.) the bands were quantitated and expressed as a ratio of GRANZYME B to TCR C α as well as a percentage relative to the CON A control. As

Fig. 4a is an autoradiograph showing the results of CAT assays of extracts prepared from U-937 cells transfected with a 1.2 kb BamHI-XhoI IL-6-CAT construct containing wild-type (WT) (lane 3, 6, 9, 12, 15, and 18). mutant M1 (lanes 2, 5, 8, 11, 14, and 17), or mutant M2 (lanes 4, 7, 10, 13, 16 and 19) kb sites as illustrated in Fig. 1. The cells were either uninduced (lanes 2 to 4) or stimulated 33 h after transfection with 10 μ g of LPS per ml (lanes 5 to 7). 2 μ g of PHA per ml (lanes 8 to 10). 100 μ g of double-stranded RNA poly(IC) (Pl-pC) per ml (lanes 14 to 16), or 100 units of IFN- γ per ml (lanes 17 to 19) for 19 h.

Fig. 4b is a graph of relative CAT activity after induction by various stimuli in U-937 cells. Acetylated and nonacetylated forms of chloramphenicol from the assay were quantitated by liquid scintillation counting. Values are represented as fold induction of the wild-type (WT) or mutant IL-6 promoter over CAT activity expressed in uninduced cells. Similar results were obtained in three independent experiments. Induction of IL-6-CAT expression in U-937 cells depends on an intact kb site.

Fig. 5 is a set of three graphs illustrating the effect of transforming growth factor (TCR) crosslinking on proliferation of the IS-2.15 clone. Each point represents the mean of triplicate determinations \pm SEM. As a positive control, the same experimental procedure was undertaken using 10^5 NOD spleen cells.

Figs. 6A and 6B are a pair of graphs illustrating that IS-2.15 supernatant specifically inhibits IL-2 dependent proliferation in IL-2 dependent murine T-cell lines. Fig. 6A shows the effect of supernatant (50% final

dilution) from 3 different clones upon T-cell proliferation. Each point represents mean \pm SEM of triplicate determinations. Fig. 6B represents the effect of titrating IS-2.15 supernatant versus different concentrations of IL-2.

5 Fig. 7 is a graph illustrating that the inhibitory activity of IS-2.15 is not reversible. Results represent cpm \pm 1 SD of triplicates.

 Fig. 8 is a graph illustrating that IS-2.15 supernatant does not modify IL-2R α expression on HT-2 cells.

10 Fig. 9 is a graph illustrating that the molecular weight of IS-2.15 supernatant suppressor activity is between 10 and 30 KD.

 Fig. 10 is a diagram of plasmid, IL-10/INSULIN-pSf3. It is 6.05 kb in size. The IL-10 gene was cloned into the plasmid, pSf3, and it is under the transcriptional control of the insulin promoter.

15 Fig. 11 is a diagram of plasmid, IL-10/IgH-pTZ19R. It is 5.82 kb in size. The IL-10 gene was cloned into the plasmid, pTZ19R, and it is transcribed under the control of the IgH enhancer/fos promoter.

 Fig. 12 is a diagram of plasmid, IL-10/IL-6P-pTZ19R. It is 6.27 kb in size. The IL-10 gene was cloned into the pTZ19R plasmid and it is transcribed under the control of the IL-6 promoter.

20 Fig. 13 is a diagram of plasmid, IL-10/HTLV1-pcD-SR α . It is 5.10 kb in size. The IL-10 gene was cloned into the pcD-SR α plasmid and it is transcribed under the control of the HTLV1 LTR.

 Fig. 14 is a diagram of plasmid, IL-10/IL-2-pTZ19R. It is 5.50 kb in size. The IL-10 gene was cloned into the pTZ19R plasmid and it is transcribed under the control of the IL-2 promoter.

Fig. 15 is a diagram of constructs expressing CTLA4-Ig under the control of various promoters.

The following will now be described in detail:

I. Inhibition of rejection of transplanted tissue and prevention of tissue destruction due to autoimmune disease and II. Isolation and characterization of the novel suppressor factor.

I. Inhibition of Rejection of Transplanted Tissue and Prevention of Tissue Destruction Due to Autoimmune Disease

There follows a detailed description of the methods by which rejection of organ and tissues and prevention of tissue destruction due to autoimmune disease can be inhibited according to methods of the invention.

A central feature of the invention is the design of therapies which specifically suppress the local immune response at the site of potential rejection or autoimmune destruction of the tissue or organ. This strategy, which targets immunosuppression to the specific anatomical region where it is needed, represents an improvement over existing clinical therapies, all of which involve general suppression of the immune response. To utilize the methods for treatment of the varied conditions described herein modifications of the general technique may be employed.

The protein encoded by the DNA transfecting cells of the transplanted tissue can exert its immunosuppressive effects in any of several ways, for example: (1) the protein itself can exert a direct immunosuppressive effect; (2) the protein can be an enzyme which acts on a substrate to convert a polypeptide substrate into an active immunosuppressive compound; or (3) the protein can be an enzyme which is capable of activating a prodrug which, when activated, is an immunosuppressive compound.

In the aspect of the invention which features prevention of transplanted tissue or organ rejection, the recombinant gene is, preferably, introduced into the cells of the tissue/organ prior to transplantation of the tissue/organ into the mammal. This can be accomplished by ex vivo transfection of isolated cells of the tissue/organ, or by transfection of the tissue/organ itself. Introduction of the DNA can also be accomplished by transfection of the tissue/organ in vivo, for example, by introducing suitably prepared DNA directly into the tissue/organ of the mammal following transplantation. Another method for the introduction of DNA into an organ involves the generation of transgenic animals in which the appropriate genes are expressed in the organ to be transplanted.

In an alternative method, the transfected cells may be in the area surrounding the transplant, but not a part of the actual transplant. For example, cultured myoblasts or immortalized renal epithelial cells may be transplanted. The transfected cells may then be intermixed with islet allografts, or may be injected into or near organ allografts (i.e. under the renal capsule).

Following transplantation, the proteins are expressed and act to inhibit rejection of the transplanted tissue/organ.

In the aspect of the invention which features the treatment of autoimmune disease, the recombinant gene is introduced into the affected tissues or surrounding tissues, preferably prior to localized onset of the disease phenotype. For example, in patients suffering from systemic lupus erythematosus (SLE), the recombinant gene may be introduced into the kidney or cardiac muscle tissues. In prediabetic patients it is preferable to introduce the recombinant transgene into the β -cells of the pancreas. The

preferred target tissues for treating rheumatoid arthritis patients are the synovial tissues of the joints, and the brain and myelin sheath are preferred targets in patients suffering from multiple sclerosis. As with therapies for transplantation, cells such as myoblasts and renal epithelia cells may be transplanted into the effected area after they are altered to express the transgene appropriately.

Construction of Recombinant Genes

Recombinant proteins capable of mediating inhibition of rejection of a transplanted tissue/organ by modulating the localized immune response include, but are not limited to, naturally occurring proteins such as IL-10, TGF- β and other IL-2 suppressor factors such as that produced by the T cell clone, IS-2.15 (described more fully below). In addition, DNA encoding non-naturally occurring polypeptides such as IL2Ig, CD2Ig, and CTLA-4 Ig fusion proteins may also be employed (see below). The genes encoding IL-10 and TGF- β have been cloned into plasmid vectors as described below. The expression of these genes can be driven by any one of a number of promoters which are described in Table 1 and Table 2.

TABLE 1

PROMOTER CONSTRUCTS

PROMOTER TYPE	PROMOTER ELEMENT	REFERENCES
CONSTITUTIVE		
	β -actin	Liu et al., <i>Mol. Cell Biol.</i> 10:3432-40 (1990)
	tubulin	Angelichio et al., <i>Nucleic Acids Res.</i> 19:5037-43 (1991)
	CMV	see Invitrogen
	SV40 enhancer	see Pharmacia
	RSV-LTR	see Invitrogen
	Adenovirus enhancer	Inoue et al., <i>Biochem Biophys Res Commun</i> 173:1311-6 (1990)
TISSUE-SPECIFIC		
	LIVER	
	serum amyloid A	Li et al., <i>Nucleic Acids Res</i> 20:4765-72 (1992)
	phenylalanine	Wang et al., <i>J Biol Chem</i> 269:9137-46 (1994)
	hydroxylase	
	IGFBP-1	Babajko et al., <i>PNAS</i> 90:272-6 (1993)
	apolipoprotein B	Brooks et al., <i>Mol Cell Biol</i> 14:2243-56 (1994)
	albumin	Pinkert et al., <i>Genes Dev</i> 1:268-76 (1987)
	vitellogenin	Corthesy et al., <i>Mol Endocrinol</i> 5:159-69 (1991)
	angiotensinogen	Brasier et al., <i>Embo J</i> 9:3933-44 (1990)
	haptoglobin	Yang et al., <i>Genomics</i> 18:374-80 (1993)
	PEPCK	Short et al., <i>Mol Cell Biol</i> 12:1007-20 (1992)
	factor IX	Jallat et al., <i>Embo J</i> 9:3295-301 (1990)

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TABLE 1 - Continued

PROMOTER TYPE	PROMOTER ELEMENT	REFERENCES
	transferrin	Idzerda et al., <i>Mol Cell Biol</i> 9:5154-62 (1989)
	β -fibrinogen	Dalmon et al., <i>Mol Cell Biol</i> 13:1183-93 (1993)
	kininogen	Chen et al., <i>Mol Cell Biol</i> 13:6766-77 (1993)
	CRP	Toniatti et al., <i>Mol Biol Med</i> 7:199-212 (1990)
KIDNEY		
	renin	Fukamizu et al., <i>Biochem Biophys Res Commun</i> 199:183-90 (1994)
HEART		
	cardiac myosin light	Lee et al., <i>J Biol Chem</i> 267:15875-85 (1992)
	chain	
	cardiac troponin C	Parmacek et al., <i>Mol Cell Biol</i> 12:1967-76 (1992)
	α -cardiac myosin	Gulick et al., <i>J Biol Chem</i> 266:9180-5 (1991)
	heavy chain	
	MCK	Johnson et al., <i>Mol Cell Biol</i> 9:3393-9 (1989)
	troponin I	
	atrial natriuretic factor	Rockman et al., <i>PNAS</i> 88:8277-81 (1991) erratum 88(21):9907
LUNG		
	pulmonary surfactant	Glasser et al., <i>Am J Physiol</i> L349-56 (1991)
	protein SP-C	
PANCREAS/ISLET		

TABLE 1 - Continued

PROMOTER TYPE	PROMOTER ELEMENT	REFERENCES
	insulin	Dandoy et al., <i>Nucleic Acids Res</i> 19:4925-30 (1991); and Selden et al., <i>Nature</i> 321:525-8 (1986)
	pancreatic amylase	Osborn et al., <i>Mol Cell Biol</i> 7:326-34 (1987)
BRAIN/GLIA		
	GFAP	Brenner et al., <i>J Neurosci</i> 10:30-7 (1994)
	JCV	Henson et al., <i>J Biol Chem</i> 269:1046-50 (1994)
	MBP	Miskimins et al., <i>Brain Res Dev Brain Res</i> 65:217-21 (1992)
	serotonin 2 receptor	Ding et al., <i>Brain Res Mol Brain Res</i> 20:181-91 (1993)
	myelin PO	Monuki et al., <i>Mech Dev</i> 42:15-32 (1993)
	myelin proteolipid	Berndt et al., <i>J Biol Chem</i> 267:14730-7 (1992)
	protein	
INDUCIBLE		
A) IMMUNE RESPONSE/NATURAL		
	IL-2	Thompson et al., <i>Mol Cell Biol</i> 12:1043-53 (1992)
	IL-4	Todd et al., <i>J Exp Med</i> 177:1663-74 (1993)
	IL-6	Libermann et al., <i>Mol Cell Biol</i> 10:2327-34 (1990); and Matsusaka et al., <i>PNAS</i> 90:10193-7 (1993)
	IL-8	Matsusaka et al., <i>PNAS</i> 90:10193-7 (1993)
	IL-10	Kim et al., <i>J Immunol</i> 148:3618-23 (1992)
	TNF- α	Drouet et al., <i>J Immunol</i> 147:1694-700 (1991)
	IL-1	Shirakawa et al., <i>Mol Cell Biol</i> 13:1332-44 (1993)

TABLE 1 - Continued

PROMOTER TYPE	PROMOTER ELEMENT	REFERENCES
	MIP-1	Grove et al., <i>Mol Cell Biol</i> 13:5276-89 (1993)
	IFN- γ	Penix et al., <i>J Exp Med</i> 178:1483-96 (1993)
	VCAM-1	Iademarco et al., <i>J Biol Chem</i> 267:16323-9 (1992)
	ICAM-1	Voraberger et al., <i>J Immunol</i> 147:2777-86 (1991)
	ELAM-1	Whelan et al., <i>Nucleic Acids Res</i> 19:2645-53 (1991)
	tissue factor	Mackman et al., <i>J Exp Med</i> 174:1517-26 (1991)
	IFN- β	Visvanathan et al., <i>Embo J</i> 8:1129-38 (1989)
	c-jun	Muegge et al., <i>PNAS</i> 90:7054-8 (1993)
	junB	Nakajima et al., <i>Mol Cell Biol</i> 13:3017-41 (1993)
	c-fos	Morgan et al., <i>Cell Prolif</i> 25:205-15 (1992)
	iNOS	Xie et al., <i>J Exp Med</i> 177:1779-84 (1993)
	G-CSF	Shannon et al., <i>Growth Factors</i> 7:181-93 (1992)
	GM-CSF	Miyatake et al., <i>Mol Cell Biol</i> 11:5894-901 (1991)
B) IMMUNE RESPONSE/SYNTHETIC multiple copies of binding sites		
	NF-KB	Lenardo et al., <i>Cell</i> 58:227-9 (1989)
	NF-IL6	Akira et al., <i>Embo J</i> 9:1897-906 (1990)
	IL6-response element	Wegenka et al., <i>Mol Cell Biol</i> 13:276-88 (1993)
	CRE	Brindle et al., <i>Curr Opin Genet Dev</i> 2:199-204 (1992)
	AP-1	Auwerx et al., <i>Oncogene</i> 7:2271-80 (1992)

TABLE 1 - Continued

PROMOTER TYPE	PROMOTER ELEMENT	REFERENCES
	p91/stat	Larner et al., <i>Science</i> 261:1730-3 (1993)
	combinations of multiple NF-KB and NF-IL6 or combinations with the other elements	
C) EXOGENOUS/NON-MAMMALIAN		
	IPTG inducible/lac	see Stratagene LacSwitch™, La Jolla, CA
	repressor/operon system	
	ecdysone-inducible	Burtis et al., <i>Cell</i> 61:85-99 (1990)
	promoter/ecdysone receptor	
	Na-salicylate-inducible promoter	Yen, <i>J Bacteriol</i> 173:5328-35 (1991)
	PG/regulator nahR	
	nalidixic acid	Rangwala et al., <i>Biotechnology</i> 9:477-9 (1993)
	inducible recA	
	promoter	

TABLE 2

PREFERRED THERAPEUTIC PROMOTER COMBINATIONS

GRAFT REJECTION	TRANSPLANT ORGAN OR TISSUE	PROMOTER CONSTRUCTS CONSTITUTIVE	TISSUE-SPECIFIC	IMMUNE RESPONSE INDUCIBLE	SYNTHETIC INDUCIBLE	ORALLY INDUCIBLE	COMBINATIONS
	Heart	CMV	cardiac troponin C atrial natriuretic factor	interleukin-6 interleukin-8	combination of multiple copies of NF- κB, NF-IL6 and AP-1	Na-salicylate- inducible promoter PG/regulator nahR	IL-6 + additional copies of NF-κB or NF-IL6 + Na- salicylate-inducible promoter PG/regulator nahR
	Lung	CMV	pulmonary surfactant protein SP-C	interleukin-6 interleukin-8	combination of multiple copies of NF- κB, NF-IL6 and AP-1	Na-salicylate- inducible promoter PG/regulator nahR	IL-6 + additional copies of NF-κB or NF-IL6 + Na- salicylate-inducible promoter PG/regulator nahR
	Kidney	CMV	renin	interleukin-6 interleukin-8	combination of multiple copies of NF- κB, NF-IL6 and AP-1	Na-salicylate- inducible promoter PG/regulator nahR	IL-6 + additional copies of NF-κB or NF-IL6 + Na- salicylate-inducible promoter PG/regulator nahR
	Liver	CMV	CRP β-fibrinogen	interleukin-6 interleukin-8	combination of multiple copies of NF- κB, NF-IL6 and AP-1	Na-salicylate- inducible promoter PG/regulator nahR	IL-6 + additional copies of NF-κB or NF-IL6 + Na- salicylate-inducible promoter PG/regulator nahR
	Bone marrow	CMV	immunoglobulin heavy chain enhancer	interleukin-6 interleukin-8	combination of multiple copies of NF- κB, NF-IL6 and AP-1	Na-salicylate- inducible promoter PG/regulator nahR	IL-6 + additional copies of NF-κB or NF-IL6 + Na- salicylate-inducible promoter PG/regulator nahR
	Pancreas/islet	CMV	insulin	interleukin-6 interleukin-8	combination of multiple copies of NF- κB, NF-IL6 and AP-1	Na-salicylate- inducible promoter PG/regulator nahR	IL-6 + additional copies of NF-κB or NF-IL6 + Na- salicylate-inducible promoter PG/regulator nahR

AUTOIMMUNITY

DISEASE	interleukin-6 interleukin-8	combination of multiple copies of NF- κB, NF-IL6 and AP-1	Na-salicylate- inducible promoter PG/regulator nahR	IL-6 + additional copies of NF-κB or NF-IL6 + Na- salicylate-inducible promoter PG/regulator nahR	TARGET
Rheumatoid Arthritis	CMV				synovium
Multiple Sclerosis	CMV	myelin basic protein JCv glial fibrillary acidic protein	combination of multiple copies of NF- κB, NF-IL6 and AP-1	IL-6 + additional copies of NF-κB or NF-IL6 + Na- salicylate-inducible promoter PG/regulator nahR	brain
Systemic Lupus erythematosus	CMV	renin cardiac troponin C	combination of multiple copies of NF- κB, NF-IL6 and AP-1	IL-6 + additional copies of NF-κB or NF-IL6 + Na- salicylate-inducible promoter PG/regulator nahR	kidney, heart
Diabetes	CMV	insulin	combination of multiple copies of NF- κB, NF-IL6 and AP-1	IL-6 + additional copies of NF-κB or NF-IL6 + Na- salicylate-inducible promoter PG/regulator nahR	pancreas

These promoters were selected because they either control expression of the proteins in an inducible manner, are tissue specific, or cause the protein to be expressed constitutively in the localized cells into which the DNA is transfected. The recombinant technology required to clone and express such genes is standard in the art and is described in detail, for example, in Sambrook et al. (Molecular Cloning. A Laboratory Manual. Cold Spring Harbor Press, Cold Spring Harbor, NY, 1989).

Promoters. Examples of promoters conferring constitutive, immunoresponsive, tissue specific or inducible expression on genes include, but are not limited to, the promoters provided in Table 1, Table 2 and Figs. 10-15.

Regulatory elements of the IL-6 promoter. The following is a detailed discussion of the IL-6 regulatory elements. Equivalent information is widely available for other promoters described herein.

Interleukin-6 (IL-6) is a cytokine released by a variety of cells including macrophages, endothelial cells, fibroblasts, T cells, glial cells and B cells in response to a variety of stimuli (Kishimoto, 1989 Blood, 74:1-10). In particular IL-6 gene expression is stimulated by all substances that trigger an immune response and inflammation, the same agents released during acute rejection (Kishimoto, 1989 Blood, 74:1-10). The IL-6 promoter is composed of a variety of overlapping regulatory elements (Kishimoto, 1989 Blood, 74:1-10). The IL-6 promoter has been cloned and inserted upstream of the chloramphenicol acetyltransferase gene. A 1.2 kb fragment of the 5' flanking region of the IL-6 gene (Ray, et al., 1988 Proc. Natl. Acad. Sci. USA, 85:6701-6705; Yasukawa, et al., 1987 Embo J., 6:2939-2945) contains all the necessary elements for induction of the IL-

6 gene. Analysis of the promoter region of the IL-6 gene revealed the presence of a putative binding site for the transcription factor NF-kB (Libermann, et al., 1990 Mol. Cell. Biol., 10:2327-2334).

5 *Putative regulatory elements of the IL-6 promoter.*

To characterize the functional role of the putative NF-kB binding site, mutations were introduced into the IL-6kB site using synthetic oligonucleotides and the gapped/heteroduplex method (Stewart, et al., 1988 Biotechniques, 6:511-518) and
10 examined for their effects on inducibility of the IL-6 gene. Enhancer activity was measured as the ability of the different enhancer constructs to induce transcription of the chloramphenicol acetyltransferase (CAT) gene after transient expression in a variety of cell types (Libermann, et al.,
15 1990 Mol. Cell. Biol., 10:3155-3162; Gorman, et al., 1982 Mol. Cell. Biol., 2:1044-1051; Gilman, et al., 1986 Mol. Cell. Biol., 6:4305-4316). Transfections of U-937 monocytic cells, Jurkat T cells and HeLa cells were carried out using the DEAE-dextran method (Pierce, et al., 1988 Proc. Natl.
20 Acad. Sci. USA, 85:1482-1486). The apparent NF-Kb binding site is an indispensable component of the IL-6 control region. The IL-6 promoter Kb site binds NF-Kb and its alteration abolishes inducibility of chimeric genes driven by the IL-6 promoter (Fig. 4) (Libermann, et al., 1990 Mol.
25 Cell. Biol., 10:2327-2334). It appears that the ability of various agents including LPS, PMA, TNF- α and dsRNA to induce IL-6 expression is a consequence of their ability to activate NF-Kb. A second enhancer element, NF-IL-6, was also shown to respond to IL-1, IL-6, TNF α , and LPS (Isshiki,
30 et al., 1990 Mol. Cell. Biol., 10:2757-2764) and recent evidence suggests that NF-kB interacts cooperatively with the NF-IL-6 factor (Mukaida, et al., 1990 J. Biol. Chem., 265:21128-21133). Mutations in either elements will abolish

inducibility. Ray et al. identified an additional element, MRE, that overlaps with a potential cAMP responsive element (CRE) (Ray et al., 1989 Mol. Cell. Biol., 9:5537-5547). The MRE is apparently induced by phorbol ester, IL-1, TNF- α and forskolin (Ray et al., 1989 Mol. Cell. Biol., 9:5537-5547). In addition, two putative glucocorticoid response elements and an AP-1 binding site have been identified in the IL-6 promoter (Kishimoto, 1989 Blood, 74:1-10). Potential binding sites for HLH and ets related transcription factors are also present, as well as a putative binding site for factors of the GATA gene family (Martin, et al., 1990 Nature, 344:444-447). The IL-6 promoter can be used as an inducible promoter for IL-10 and TGF- β expression (see Dendorfer et al., Mol. Cell. Biol., 14:4443-4454 (1994), incorporated herein by reference).

Additional Promoters.

Publicly available promoters that are inducible by factors involved in the immune response include NF κ B and NF-IL-6, both of which are induced by IL-1 and other immune response mediators; interferon response element, which is induced by γ -interferon; IL-6 response element, which is induced by IL-6; CRE, which is induced by prostaglandins.

Promoters that are inducible by exogenous factors include the glucocorticoid response element, which is induced by glucocorticoids; retinoic acid response element, which is induced by retinoic acid; NF- κ and NF-IL-6, which are induced by inflammatory agents; Verapamil response element and OS-1 response element, which are induced by verapamil and OS-1; glucose response element, which is induced by glucose; Mouse Mammary Tumor Virus LTR, which is induced by dexamethasone.

Examples of promoters that confer constitutive expression include, but are not limited to, the following:

The human T cell leukemia virus type 1 long terminal repeat; the Rous sarcoma virus long terminal repeat; the Moloney murine leukemia virus long terminal repeat; the simian virus 40 early enhancer/promoter; the cytomegalovirus immediate early promoter; the β -actin promoter; the islet cell insulin promoter; the B cell immunoglobulin enhancer/promoter; the T cell receptor enhancer promoter that is T cell specific, such as the HTLV-1 promoter, or an endothelial cell specific promoter.

10 Plasmids or other vectors encoding most of these promoters are commercially available. Where this is not the case, the promoters can be isolated and cloned into appropriate plasmids or vectors by conventional recombinant techniques following the directions provided in Sambrook et al. (*Supra*).

Genetic Constructs.

Examples of genetic constructs which have been made are include IL-10 cDNA driven by the insulin promoter (Fig. 10); IL-10 cDNA driven by the immunoglobulin heavy chain promoter (Fig. 11); IL-10 cDNA driven by the IL-6 promoter (Fig. 12) ; IL-10 cDNA under the control of the SR α promoter (Fig. 13) (Takebe, Y. et al. 1988, Mol. Cell Biol., 8:466), and; IL-10 cDNA driven by the IL-2 promoter (Fig. 14).

Similarly, the human TGF- β 1 cDNA (Derynck, et al., 1985 Nature, 316:8) and other genes for immunosuppressive proteins can be inserted into such vectors. For inducible expression, cDNAs can be cloned directly downstream of the IL-6 (Libermann, et al., 1990 Mol. Cell. Biol., 10:2327-2334) or the IL-2 (Hoyos, et al., 1989 Science, 244:457-460; Fraser, et al., 1991 Science, 251:313-316) promoters. Both promoters are highly inducible by several agents, particularly substances released from activated macrophages

during immune response (Kishimoto, 1989 Blood, 74:1-10; Crabtree, 1982, Science, 243:355-361).

To evaluate expression of IL-10 or TGF- β by the transfected cells, several different assays, such as the IL-10 ELISA assay, described in Current Protocols in Immunology (Mosmann, 1991 In: Current Protocols in Immunology; 1:6.14.1-6.14.8) herein incorporated by reference, can be used. IL-10 activity is determined by the ability of IL-10 to inhibit cytokine production by activated macrophages and this can be measured in a bioassay (Fiorentino, D.F. 1991, J. Immunol., 147:3815), herein incorporated by reference). TGF- β expression can be measured by either the thymocyte proliferation assay in the presence or absence of neutralizing anti-TGF- β 1 antibodies (Genzyme Corporation, Cambridge, MA) or by a radioreceptor assay using A549 lung carcinoma cells (deposited with the ATCC) as described (Mosmann, 1991 In: Current Protocols in Immunology; 1:6.14.1-6.14.8).

Activation of a pro-drug

A gene encoding an enzyme that activates an immunosuppressive drug can be cloned and expressed in T-cells or in a graft. Expression of a gene such as cyclosporine synthetase (Lawen, A. 1990, J. Biol. Chem., 265:11355) by the graft would activate the drug, e.g., a cyclosporine precursor, and create an environment of local immunosuppression, thus allowing improved survival of the grafted tissue or organ.

Another immunosuppressive agent that must be metabolized before becoming active is cyclophosphamide. Cyclophosphamide metabolism normally takes place in liver microsomes. The genes encoding the activating enzymes may be cloned and expressed in T cells or engrafted organs, resulting in local activation of the drug.

Enzymes that act to produce a compound capable of inhibiting rejection of a tissue or organ.

Molecules that are the target for preformed antibodies, which present the biggest barrier for xenotransplants, are largely molecules that are heavily glycosylated (Geller et al, 1992, Transplant. Proc., 24:592). The glycosylation sites appear to be targets for these antibodies. Antigenic determinants are frequently located on N-linked substitutions. To eliminate the targets for such xenoreactive natural antibodies, a glycosidase can be expressed in the transplanted organ.

Immunosuppressive Fusion Proteins

DNA encoding immunosuppressive fusion proteins may also be employed to prevent graft rejection and autoimmune mediated tissue destruction. Such fusion genes may encode immunosuppressive amino acid sequences from for example, IL-2, CD2 or CTLA-4 linked to the Fc portion of the Ig heavy chains. The Fc portion of the Ig heavy chains is known to stabilize soluble proteins. The IL-2 Ig fusion encoded by these constructs binds to activated lymphocytes causing direct killing through the Fc of the IgG component. The CD2 and CTLA4 Fg fusions possess an Fc portion which does not activate complement or bind with FcR⁺ phagocytes, but which does confer an extended half life. Thus, the CD2 and CTLA4 fusion molecules serve to actively block costimulatory T-cell activation signals. CTLA4 Ig proteins and DNA encoding the same are described in Linsley et al., J. Exp. Med., 174:561-569 (1991); Lenschow et al., Science, 257:789-792 (1992); Linsley et al., Science, 257:792-795 (1992); and Turka et al., PNAS, 89:11102-11105 (1992) and DNA encoding the same are described in Steurer et al., J. Amer. Soc. Nephrol., 4:919 (1993) and Kato et al., J. Exp.

Med., 176:1241-1249 (1992). IL-2 immunoligands and DNA encoding the same are described in Landolfi, J. Immunol., 146:915-919 (1991), Vie et al., PNAS, 89:11337-11341 (1992), and Steele et al., J. Amer. Soc. Nephrol., 4:636. All these
5 references are incorporated herein.

Introduction of Recombinant Genes into Cells of Tissues

A number of methods can be employed for the introduction of recombinant genes into cells of a tissue or organ. In addition to those examples provided below, Table
10 3 provides further examples of methods which may be used to deliver DNA both *in vitro* and *in vitro*.

TABLE 3

EXAMPLES OF GENE TRANSFER PROTOCOLS*

TARGET TISSUE	METHOD	REFERENCES
A. General	Liposome	Nabel et al., PNAS USA, 90:11307-11 (1993) Nabel et al., Hum. Gene Ther., 3:649-56 (1992) Zhu et al., Science, 261:209-11 (1993)
	Particle Bombardment	Cheng et al., PNAS USA, 90:4455-9 (1993)
	Microcapsules	Chang et al., Hum Gene Ther., 4:433-40 (1993)
B. Muscle - generally	Retroviral	Dai et al., PNAS USA, 89:10892-5 (1992)
	Cell transplantation	Gussoni et al., Nature, 356:435-8 (1992)
	Adenoviral	Ragot et al., Nature, 361:647-50 (1993)
	Liposome	Nabel et al., J. Clin. Invest., 91:1822-9 (1993) Nabel et al., PNAS USA, 89:5157-61 (1992)
	In Vivo	Kass-Eisler et al., PNAS USA, 90:11498-502 (1993)
C. Cardiac Muscle	In Vitro	Kass-Eisler et al., PNAS USA, 90:11498-502 (1993)
	Adenovirus	Kass-Eisler et al., PNAS USA, 90:11498-502 (1993)
	Retroviral	Kay et al., Hum. Gene Ther., 3:641-7 (1992) Rettinger et al., PNAS USA, 91:1460-4 (1994) Cardoso et al., Hum. Gene Ther., 4:411-8 (1993)
D. Liver	Adenoviral Vector	Kay et al., PNAS USA, 91:2353-7 (1994)
	Herpes Virus	Andersen et al., Cell Mol. Neurobiol, 13:503-15 (1993)
	Liposomes	Jiao et al., Exp. Neurol., 115:400-13 (1992)

TARGET TISSUE	METHOD	REFERENCES
F. Neurons & Ganglia	Adenoviral	La Salle et al., Science, 259:988-90 (1993)
G. Glial Cells	Glioma	Yamada et al., Jpn J Cancer Res., 83:1244-7 (1992)
	Retroviral	Yamada et al., Jpn J Cancer Res., 83:1244-7 (1992)
H. Lung	Adenovirus	Flotte et al., PNAS USA, 90:10613-7 (1993)

Transfection of Dispersed Cells from a Tissue or Organ

Transfection of dispersed cells from a tissue/organ can be accomplished by any number of known methods, including calcium phosphate precipitation, DEAE dextran, electroporation, and liposome mediated transfection. The method to be used for any given population of cells will depend upon the cell type that is being transfected and their sensitivity to the components of the transfection mixture. Each of these methods is well known to those skilled in the art, and are described in detail.

Viral Delivery. Retroviral vectors, or other viral vectors, such as adenoviral vectors, with the appropriate tropisms for cells useful for therapeutic delivery, may be used as a gene transfer delivery system for the methods of the invention. Numerous vectors useful for this purpose are generally known have been described (Miller, Human Gene Therapy, pp. 15-14 (1990); Friedman, Science, 244:1275-1281 (1989); Eglitis and Anderson, BioTechniques, 6:608-614 (1988); Tolstoshev and Anderson, Current Opinion in Biotechnology, 1:55-61 (1990); Sharp, The Lancet, 337:1277-1278 (1991); Cornetta et al., Nucleic Acid Research and Molecular Biology, 36:311-322 (1987); Anderson, Science, 226:401-409 (1984); Moen, Blood Cells, 17:407-416 (1991); and Miller and Rosman, Biotechniques, 7:980-990 (1989)). Retroviral vectors are particularly well developed and have been used in a clinical setting (Rosenberg, et al., N. Engl. J. Med., 323:370 (1990)).

The retroviral constructs, packaging cell lines and delivery systems which may be useful for this purpose include, but are not limited to, one, or a combination of, the following: Moloney murine leukemia viral vector types; self inactivating vectors; double copy vectors; selection

marker vectors; and suicide mechanism vectors. The Moloney murine leukemia retroviral system of IP-10 delivery is particularly useful since it targets delivery of the IP-10 protein to the hematopoietic cells which may be used for autologous or non-autologous therapy.

Non viral methods for delivery. Nucleic acid encoding the immunosuppressant polypeptides of the invention under the regulation of the appropriate promotor, and including the appropriate sequences required for insertion into genomic DNA of the patient, or autonomous replication, may be administered to the patient using the following gene transfer techniques: microinjection (Wolff et al., Science, 247:1465 (1990)); calcium phosphate transfer (Graham and Van der Eb, Virology, 52:456 (1973); Wigler et al., Cell, 14:725 (1978); Felgner et al., Proc. Natl. Acad. Sci. USA, 84:7413 (1987)); lipofection (Felgner et al., Proc. Natl. Acad. Sci. USA, 84:7413 (1987); Ono et al., Neuroscience Lett., 117:259 (1990); Brigham et al., Am. J. Med. Sci., 298:278 (1989); Staubinger and Papahadjopoulos, Meth. Enz., 101:512 (1983)); asialorosonucoid-polylysine conjugation (Wu and Wu, J. Biol. Chem., 263:14621 (1988); Wu et al., J. Biol. Chem., 264:16985 (1989)); and electroporation (Neumann et al., EMBO J., 7:841 (1980)). These references are hereby incorporated by reference, as are the additional reference.

Transfection of Whole Tissues or Organs Tissue or organs, such as pancreatic islets, can be transfected using either electroporation (BioRad) as described by Welsh et al (Welsh, et al., 1990 Proc. Natl. Acad. Sci USA, 87:5807-5811), lipofectin reagent (GIBCO-BRL), pH-sensitive liposomes as described by Welsh et al. (Welsh, et al., 1990 Biomed. Biochim. Acta, 49:1157-1164) or microparticle bombardment with the biolistic PDS-1000/He system (BioRad)

(Yang, et al., 1990 Proc. Natl. Acad. Sci. USA, 87:9568-9572).

Transgenic Organs. Xenotransplantation of organs across species barriers has been proposed as a source of donor organs (Auchincloss, 1990 Transplant Rev., 4:14). In particular, experiments with pigs as xenograft donors for human patients are in progress (Gustafsson, et al., 1990 J. Immunol., 145:1946-1951). To prevent xenograft rejection, IL-10, TGF- β , or another protein with immunosuppressive activity can be expressed in the transgenic organ in either a tissue-specific or inducer-specific manner. In order to express IL-10 and/or TGF- β in either pancreatic islets or T cells of transgenic animals, the expression vector constructs described above can be used. For example, the HTLV I LTR driven IL-10 and TGF- β constructs can be used for T cell-specific expression, and the insulin promoter-driven IL-10 and TGF- β constructs can be used for pancreatic islet cell-specific expression.

For production of transgenic animals, expression vector constructs will be linearized and injected into oocytes according to standard protocols (Sarvetnick, et al., 1988 Cell, 52:773-782). Expression of the transgenes can be evaluated by sacrificing a second generation animal and isolating MRNA from different organs and from multiple hematopoietic cell types. Since constitutive expression of such genes in transgenic animals might lead to severe developmental defects or other pathological changes, the transgenic animals preferably contain genes of interest under the transcriptional control of inducible promoters as described above, such as the IL-6 promoter. IL-6 promoter driven constructs should be inducible in a variety of different cell types including endothelial cells, T cells,

macrophages, fibroblasts, stromal cells, and mesangial cells (Kishimoto, 1989 Blood, 74:1-10).

Evaluation of Constructs.

DNA constructs employed in the invention may be tested *in vitro* for their general immunosuppressive effect *in vitro* using the MCR assay (Coligan et al., *Current Protocols in Immunology* (1992) Wiley & Sons, Eds.). Those constructs which confer a [20%] or greater decrease in proliferation relative to the control are useful in the methods of the invention.

Constructs may also be readily tested *in vivo* using the mouse islet transplant assay (see below) or other transplant and autoimmune modes readily available to those in the field. Promoters which may be used for the expression of the immunosuppressive proteins or glycosidases of the invention are shown in TABLE 1. Preferred promoter-gene combinations for the treatment of a graft rejection of various tissues are shown in TABLE 2, top and preferred promoter-gene constructs for the treatment of specific autoimmune diseases are shown in TABLE 2, bottom.

The methods of the invention may be used alone for the treatment of autoimmune disease or for the prevention of graft rejection or they may be used in combination with other known local or general immunosuppressive therapies.

Constructs of the invention which are immunosuppressive, as used herein, are those constructs which prolong engraftment beyond the 50% rejection period for the control animal similarly treated, but lacking the transgene. For example, constructs which include non-specific, constitutative, immune-responsive, inducible promoters, or β -cell specific promoters may be tested in the pancreatic islet cell transplant model described herein. Tissue specific expression may be listed in the appropriate

cell type, or in transgenic mice, as discussed above. Constructs for use in the treatment of autoimmune disease may be evaluated by examination of tissue injury in the appropriate model. For example, mice with the epr gene, NOD mice and EAE mice may be utilized. In epr mice tissue injury in the kidney may be evaluated by immunopathologic techniques. For example, MRL-Lpr mice receiving epithelial cells which produce GM-CSF under the renal capsule may be compared to genetically identical mice receiving epithelial cells not containing the GMZSF encoding transgene. In such a test, those constructs which confer increased tissue survival relative to the untreated control are useful in the methods described herein. Useful constructs are also those which provide a mixed lymphocyte reaction (MLR) by decreasing proliferation by 20%, more preferably 40% and most preferably by 60% relative to control cells not expressing the transgene. This assay may be done as described in *Current Protocols in Immunology* (1992, Wiley & Sons, Eds., Coligan et al.).

A. Novel Suppressor Factor.

There follows a detailed description of the transplant model in which the immunosuppressive effects of the suppressor factor of the invention were observed, followed by a description of the cells from which the factor was isolated, the procedure used for isolation, and the experiments in which immunosuppression was observed.

Use. Uses of the suppressor factor protein of the invention are based on its unique properties, in particular, its ability to block IL-2 stimulated T-cell proliferation without being cytotoxic or inhibiting the expression of IL-2 or IL-2 receptor. Uses fall into two main categories: diagnosis and therapy.

A nucleic acid encoding the suppressor factor of the invention can be used therapeutically to alter the effects of IL-2 on IL-2R bearing cells, or of IL-4 on IL-4R bearing cells. Four examples of how this method could be used

- 5 include, but are not limited to, the following. 1) Cells that are targets of autoreactive IL-2R- or IL-4R- bearing cells can be transfected with a nucleic acid encoding the suppressor factor. The cell expresses and secretes the suppressor factor, which causes an alteration of the IL-2
- 10 effect, such that IL-2 dependent proliferation of autoreactive T-cells proximal to the cell are subject to inhibition by the suppressor factor. 2) Cells that are in close proximity to cells, e.g., autoreactive T-cells, as they circulate, such as the endothelial cells lining the
- 15 blood vessels, can be transfected with nucleic acid encoding the suppressor factor. The suppressor factor protein would be released into the blood stream where it alters the effect of IL-2 on circulating IL-2R bearing cells. 3) In a similar manner, epithelial cells transfected with a nucleic
- 20 acid of the invention may secrete suppressor factor protein, altering the effect of IL-2 on proximal IL-2R bearing cells. Epithelial cells, e.g., epithelial cells of the kidney proximal tubule and of the gut, activate or inactivate T-cell proliferation and are thus expected to come into
- 25 proximity with them. 4) Finally, autoreactive T-cells may themselves be transfected with the nucleic acid of the invention, such that the T-cells make and secrete suppressor factor which would in turn alter the effect of IL-2 on themselves. Cells transfected in the manner of the methods
- 30 above are expected to equally alter the effect of IL-4 on IL-4R bearing cells.

Cells to be transfected can be accessed in several ways, either by removing the cells from a patient for

transfection, or by administering a vector, e.g., a virus, comprising a suppressor factor-encoding nucleic acid to the patient. Alternatively, cells, e.g., cells of a cell line, or of a graft, can be transfected *in vitro* and then
5 introduced to the patient.

Diagnosis. Human patients suffering from autoimmune diseases will be expected to exhibit abnormally low levels of the suppressor factor of the invention in the blood and in the endocrine portion of the pancreas. Assaying blood,
10 other biological fluid samples, or pancreatic tissue samples for the factor can thus provide a means of monitoring therapy and assessing the immune status of patients with autoimmune diseases, e.g., IDDM. Such assays can be carried out by conventional immunoassay methods employing antibodies
15 to the suppressor factor, made by conventional techniques in which a rabbit is immunized with the factor and the resulting antibody harvested and labeled, e.g., with a radioisotope or a fluorescent tag.

Therapy. For use in therapy, the suppressor factor
20 protein of the invention is first purified to homogeneity using conventional techniques. It is then mixed with a pharmaceutical carrier and administered intravenously or by some other standard method, or used to perfuse a graft to be transplanted. The factor can be used to treat autoimmune
25 diseases such as systemic lupus erythematosus (SLE), type 1 diabetes, and rheumatoid arthritis, as well as other disease states involving the immune system such as graft versus host disease. Dosage will be in the range of dosages used for currently available immunosuppressive agents.

30 Cells. The anergic T-cell clone IS-2.15 is a recombinant human IL-2 (rIL-2) dependent, non-cytolytic, CD8+, V β 11+ clone propagated from the islets of 2 month old euglycemic male NOD mice.

The T-cell clone, IS-2.15, was deposited with the American Type Culture Collection on _____, and bears the accession number ATCC no. _____.

Applicants acknowledge their responsibility to replace the
5 plasmid loose viability before the end of the term of a patent issued hereon, and their responsibility to notify the American Type Culture Collection of the issuance of such a patent, at which time the deposit will be made available to the public. Prior to that time the deposit will be made
10 available to the Commissioner of Patents under the terms of CFR §1.14 and 35 USC §112.

The following examples are meant to illustrate the invention and are not meant to limit the invention.

Example 1

15 This example illustrates a method for gene transfer which may be used to administer immunosuppressive proteins to the kidney area and provides a test for the effectiveness of constructs.

Macrophage (M ϕ) colony-stimulating factor (CSF-1)
20 expression precedes renal injury in autoimmune MRL-*lpr* mice and is responsible for M ϕ proliferation and survival in glomeruli. [The purpose of this study was to test the capacity of local and systemic expression of M ϕ growth factors, CSF-1 and granulocyte-M ϕ CSF (GM-CSF), to initiate
25 renal injury in normal (C3H-++, MRL-++) and autoimmune prone (C3H-*lpr*, MRL-*lpr*), mice.] We designed a gene transfer system to deliver cytokines into the kidney by transducing TEC using retroviral vector (MFG) expressing CSF-1, GM-CSF or interleukin-6 (IL-6). Transduced cells (90%)
30 constitutively secreted high levels of stable, bioactive protein. We placed transduced primary tubular epithelial cells (TEC, 5x10⁶ cells) under the left renal capsule of

normal and autoimmune prone mice prior to renal injury and evaluated renal pathology 3, 7, 14 and 28d post-implant.

CSF-1 & GM-CSF, but not IL-6 nor uninfected TEC, caused an influx of cells intrarenally radiating from the implant site by 3-8d (mostly M ϕ) which was maximal by 15-28d (mostly T cells) in strains with *lpr*. Renal injury was most severe in the MRL-*lpr* strain. By comparison, the kidney of MRL-++ and C3H mice remained normal. Although implanted transduced TEC caused a sustained increase of CSF-1, GM-CSF and IL-6 in the circulation, the contralateral kidney remained unaffected. In conclusion, gene transfer of M ϕ growth factors into the kidney initiates severe local renal injury in autoimmune prone mice with *lpr* gene but does not compromise the kidney in non-autoimmune hosts. In addition, systemic delivery of growth factors at these concentrations does not induce renal damage. These studies offer a novel gene transfer approach to explore the impact of local and systemic cytokine production on renal injury.

TABLE 4

Donor Recipient		Intrarenal			
(strain)	(strain)	CSF-1	GM-CSF	IL-6	uninfected
C3H-++	C3H-++	0 \pm 0	0 \pm 0	0 \pm 0	0 \pm 0
C3H-++	C3H- <i>lpr</i>	1.0 \pm 0.2	1.0 \pm 0	0 \pm 0	0 \pm 0
MRL- <i>lpr</i>	MRL-++	0 \pm 0	0 \pm 0	0 \pm 0	0 \pm 0
MRL- <i>lpr</i>	MRL- <i>lpr</i>	1.6 \pm 0.2*	2.8 \pm 0.3*	0 \pm 0	0 \pm 0

n = 2-3

28d after implant

*p<0.05

Grade: 0 (none) - 3 (severe)

Example 2

This example illustrates retroviral delivery to the kidney.

We constructed a gene transfer system to deliver
5 cytokines into the kidney and circulation. We selected a
recombinant Moloney murine leukemia virus retroviral vector
(MFG) system which efficiently transfers cytokine genes
without the need of selection. Primary tubular epithelial
cells (TEC) derived from normal (C3H/Fej, MRL-++) and
10 autoimmune (C3H-*lpr*) mouse strains were cultured and
transduced with retroviral vectors expressing colony
stimulating factor-1 (CSF-1), granulocyte-macrophage CSF
(GM-CSF) and interleukin-6 (IL-6). Each cytokine was
constitutively released by most cells (90%) as determined by
15 colony forming assays and immunoperoxidase methods. Uniform
levels of cytokines were released by TEC from each strain.
These TECs remained capable of dividing and secreting stable
amounts of biologically active cytokines for >8mo in
culture. We placed transduced TEC (5×10^6) under the renal
20 capsule and monitored cytokine secretion. TEC formed nests
of cells that were vascularized by extensions from
peripheral capillaries. Co-expression of cytokine and the
 β -galactosidase gene established that TEC remained viable
and capable of producing cytokines in the kidney for at
25 least 28d. These retrovirally infected TEC delivered a
continuous supply of CSF-1, GM-CSF or IL-6 into the
circulation from 8-28d.

This example demonstrates a gene transfer system to
deliver a continuous supply of cytokines in the kidney and
30 circulation. This system offers a novel, powerful approach
for establishing the impact of sustained cytokine expression
on renal injury.

TABLE 5

5	strain *(colony forming units)	cytokine	pre-implant (CFU*)	post-implant (CFU*)	
				8d	28d
	C3H/Fej	CSF-1	0±0	1±0	12±6
	C3H- <i>lpr</i>	CSF-1	0±0	14±3	7±2
	MRL- <i>lpr</i>	CSF-1	4±0	10±3	25±5
	MRL- <i>lpr</i>	GM-CSF	0±0	13±5	15±10

10 Example 3.

This example illustrates the Allogeneic transplant model which may be used to test the effectiveness of constructs. We have investigated murine islet cell transplants as a model to study allograft rejection (Pankewycz, et al., 1989 Transplantation, 47:318; Mackie, et al., 1990 Transplantation, 49:1150) and performed allogeneic pancreatic islet cell transplants according to described techniques, herein incorporated by reference (Gotoh, et al., 1985 Transplantation, 40:437). Pancreatic islets harvested from DBA/2 (H-2^d) mice are transplanted under the renal capsule of 8-10 wk old B6AF₁ (H-2^{b/k, d}) mice previously rendered diabetic by the beta cell toxin, streptozotocin (250mg/kg ip). Following successful transplantation, blood glucose returns to normal (<200mg/dl) within 7 days. Blood glucose levels are assessed every other day throughout the first 50 days post-transplantation, then twice weekly through the next 50 days. Rejection is defined as hyperglycemia in excess of 300 mg/dl or 3 consecutive days of blood glucose >250mg/dl. This assay may be used to test immunosuppressive constructs for use in both autoimmune and transplant related therapies.

Unmodified graft rejection. Animals are sacrificed at days 4, 8, or 12. Histological examination of day 4 pancreatic islet allografts showed early leukocyte infiltration but well preserved islet morphology. Day 8 allografts showed a massive leukocyte infiltrate with severe islet injury; however, the islets were still discernable. By day 12 the islets were completely destroyed. There was early organization of the cellular exudate, and the leukocyte infiltrate was less intense than observed on day 8. Immunoperoxidase staining of day 12 allograft specimens showed that most infiltrating leukocytes were macrophages, and expressed endogenous peroxidase.

Tolerance induction. Bluestone and his colleagues first reported that high doses of hamster and anti-mouse CD3 mAb caused profound immunosuppression (Hirsch, et al, 1988 J. Immunol., 140:3766). It was noted that lower doses, closer to clinically applied doses, produced tolerance in many, but not all, islet allograft recipients (Mackie, et al., 1990 Transplantation, 49:1150). In a new study using the same high total dose employed by Bluestone but administered over a 3d period, permanent engraftment of 90% of DBA/2 into B6AF₁ islet graft recipients was obtained. Daily treatment with a recombinant IL-2 diphtheria toxin related fusion protein (DAB₄₈₆-IL-2) also causes a 50% incidence of graft tolerance (Pankewycz, et al., 1989 Transplantation, 47:318). Twice daily therapy results in 90% incidence of tolerance.

Example 4.

PCR based identification of activation associated transcripts. PCR can be used to identify activation associated transcripts. The PCR technique was adapted for mRNA phenotyping from tissues that were snap frozen in liquid nitrogen in the following fashion: (a) total

cellular RNA was isolated using the cesium chloride modification of the guanidine isothiocyanate method, (b) first-strand cDNA was synthesized using oligo (dT) primer and M-MLV reverse transcriptase, and (c) the cDNA was amplified (25 cycles) with sequence specific oligonucleotide primer-pairs and thermostable DNA polymerase (Taq DNA polymerase), using a DNA thermal cycler. Each sample as co-amplified with a β -actin oligonucleotide primer as an internal control. The PCR products were analyzed by agarose gel electrophoresis for the predicted fragment size and for specificity by Southern hybridization using ^{32}P randomly primed partial length DNA probes. In each case, the radiolabelled cDNA probes identified a single characteristic band when applied to RNA-revived from ConA + RNA activated splenic sutures. We have determined the number of cycles required for attainment of the linear portion of the radioactivity curve following hybridization with radiolabeled probes. For detection of each of the mRNAs, 25 cycles of PCR amplification was adequate for this purpose. The antisense and sense primers hybridize to sequences encoded by different exons (Table 6). This precaution reduces the possibility that DNA, rather than RNA, sequences are unwittingly selected for amplification and confused with RNA based amplification. In each experiment, one control sample lacks reverse transcriptase in order to identify preparations with contaminant DNA. The relative abundance of each amplified product was semiquantitatively evaluated using the PHOSPHOR IMAGER system (Molecular Dynamics, Inc.).

Example 5.

Correlation of allograft transcriptional activity with histology. RNA was isolated from allografts, lymphoid tissue and non-lymphoid tissue at days 1, 4 (the time of maximal cellular infiltration), 8, 12 (the time of complete

graft destruction) and day 19. Peak IL-2 and IFN γ gene transcription in the allograft closely correlated in time with the peak of leukocytic infiltration (Figs. 1, 2 and Table 7). IL-2 and IFN γ transcripts were expressed on day 8, but not at day 1, and only sporadically at days 4 and 12. Whereas IL-2 transcripts were detected in all rejecting allografts at day 8 post-transplant, IL-4 transcripts, although present in some grafts, were barely detectable at any time point. Granzyme B transcripts were detected in most allografts even as early as day 1 in some grafts. Nonetheless, co-amplification of granzyme B and TCR α sequences revealed that the ratio of granzyme B to TCR α rose progressively through days 1 to 12 (Fig. 3). The ratios of granzymes B to TCR α transcripts were compared to those obtained from a 72h Con A activated B6AF₁ spleen cell culture. β -actin transcripts were detected in all specimens. Table 7 shows the pattern of IL-2, IFN γ , IL-4 and grandzyme B transcription at Day 8 in acutely rejecting mice. It is noteworthy that detection of IL-2 and granzyme B transcripts in the allograft foreshadowed their appearance in the lymph nodes and spleen. However, the data also indicates that the pattern of transcriptional activity in the spleen at day 8 closely reflects the pattern of transcriptions in the allograft.

Example 6.

Molecular characterization of allograft tolerance.

Animals are treated with immunosuppressive regimes (e.g., anti CD3, DAB₄₈₆-IL-2, or IL-2 Fc) to induce permanent engraftment ["tolerance"] and the transcriptional activity is compared with activity in unmodified graft rejection and syngeneic grafts. Animals are sacrificed at sequential time points during the induction and maintenance phase of graft

tolerance and mRNA isolated from the graft, spleen and lymph node.

Example 7.

Transcriptional activity in unmodified, anti-CD3 or
5 DAB₄₈₆-IL-2 treated hosts. A reduction in proportion of
grafts/spleen bearing detectable IL-2 and IFN γ mRNA was
evident in hosts receiving tolerogenic therapies (Table 7).
These data suggest that TH1 subset activation is interrupted
by tolerogenic therapies. The proportion of tissues bearing
10 detectable granzyme B and IL-4 transcripts was not altered
by treatment. Our preliminary results indicate that IL-10
transcription is evident in all grafts from unmodified and
treated hosts.

The data and in the preceding two sections indicate
15 that inactivation/destruction of TH1 cells may be closely
linked to tolerance induction, and the balance between
expression of pro-inflammatory (IL-2, IFN γ) and anti-
inflammatory transcripts (IL-10) is markedly altered in
hosts receiving tolerogenic therapy.

20 Example 8.

Pathology and transcriptional events in pancreatic
islet cell allografts during rejection and tolerance
induction. Using a model of mouse pancreatic islet cell
transplantation and the RVT-PCR technique, intragraft and
25 splenic expression of IL-2, IFN γ , IL-2R α , IL-4, and
granzyme B mRNA was studied. An analysis of IL-10 mRNA was
initiated. In a totally MHC mismatched transplant (DBA2/J-
B6AF1/J), the graft is completely rejected by day 19. Mice
were transplanted and sacrificed at 1, 4, 8, 12 and 19 days
30 post transplant. Maximal mononuclear cell infiltration of
the allografts occurred at day 8. The peak expression of
IL-2 and IFN γ mRNA also occurred at day 8 while meager IL-4
transcripts were detected in some grafts at this point.

Peak expression of granzyme B occurred at days 8-12. In a second set of experiments, transplanted mice were treated either with anti-CD3 mAb or an IL-2 toxin fusion protein (DAB/486-IL-2) with doses that we have shown to significantly delay rejection and to induce tolerance in at least 90% of mice. Treated mice were sacrificed 8 days post transplant for RNA extraction. IL-2 and IFN γ transcription was markedly reduced. Preliminary data indicate that tolerogenic therapies do not block IL-10 expression.

10 Example 9.

Islet cell graft infiltrating T-cell lines and T-cell clones. Using the methods described by Fitch and Gajewski (Fitch, 1991 Current Protocols in Immunology. 3.13.1-3.13.11), both rIL-2 (80 U/ml) and rIFN γ (4000 U/ml) or a mixture of rIL-2 and Con A supernatants to derive outgrowths of islet graft infiltrating T cells were used. The IL-2 and IFN γ mixture favors TH1 cell growth while the latter mixture favors TH2 cell growth (Fitch, 1991 Current Protocols in Immunology. 3.13.1-3.13.11). The outgrowths are then cultivated in media that favors the maintenance of TH1 or TH2 cells. Other clones are derived using only rIL-2 and donor strain islets. To date we have derived at least two T-cell lines cultivated under TH1- and one cultivated under TH2- promoting growth media from day 2, 8, 12, 19 unmodified and DAB\486-IL-2 treated hosts. Aliquots of these lines have been frozen and stored. A limiting dilution method (MacDonald, et al., 1990 Immunol. Rev. 51:93) to derive T-cell graft infiltrating clones from day 8 unmodified and DAB\486-IL-2 treated hosts was used. The media used to selectively propagate TH1 and TH2 clones does yield a high proportion of CD4+ clones while use of a protocol utilizing IL-2 and islet cell cultures primarily yields CD8+ T-cells.

Example 10.

Suppressive effects of IS-2.15 T-cells. Transfer of IS-2.15 T-cells into prediabetic NODs prevents rapid induction of diabetes of accelerated diabetogenic autoimmunity (Pankewycz, O., et al., 1991, Eur. J. Immunol., 21:873). The suppressive effects of IS-2.15 are characterized in the following detailed description.

H-4, G-3, #3 and #5 cells are CD3+, CD8+ islet-infiltrating T-cell clones isolated from the islets of 2 month old euglycemic NOD mice as previously described (Pankewycz, O., et al., 1991, Eur. J. Immunol., 21:873). HT-2 (Watson, J., 1979, J. Exp. Med., 150:151) and CTLL-2 (Gillis, S., et al., 1977, Nature, 268:154.) murine IL-2/IL-4 dependent T-cells were obtained from Dr. D. Perkins (Brigham and Women's Hospital, Boston, MA) and the American Type Culture Collection (Rockville, MD), respectively.

T-cells were cultured in RPMI-1640 media (Gibco, Grand Island, NY) supplemented with 10% (vol/vol) fetal bovine serum, 100µg/ml streptomycin, 100 u/ml penicillin, 10 µM HEPES, 1 µM sodium pyruvate, 10^{-5} M 2-mercaptoethanol and 1% minimal essential amino acids (complete RPMI medium). In some cultures irradiated (3,000 rads) syngeneic splenic cells were plated with IS-2.15, G-3 and H-4 cloned T-cells. T-cell growth factor (TCGF) was added to some HT-2 and CTLL-2 cell cultures.

Supernatant from NOD T-cell clones

T-cell clones were isolated from feeder cells by Lympholyte M (Cedarlane Lab. Hornby, Ontario, Ca.) gradient separation and placed in culture at a concentration of 10^6 cells/ml in complete RPMI medium for 24 hours. After this period, T-cells were pelleted by centrifugation, and the supernatant was aliquoted in siliconized tubes and stored at -70°C.

*Proliferative responses against immobilized
mitogenic mAbs*

Fifty $\times 10^3$ cloned T-cells were cultured in round
bottom 96-well plates in 0.2 ml of complete RPMI medium
5 alone or with 5×10^5 irradiated (3,000 rads) syngeneic
spleen cells and different concentrations of immobilized
anti-CD3 or anti-V β 11 mAbs. Wells were coated with mAbs for
2 hours at 37°C. Cells were cultured in the mAb coated
plates for 48 hours, pulsed with 1 μ Ci [3 H]thymidine during
10 the last 6 hours of culture, harvested with a semiautomated
cell harvester (PHD, Cambridge, MA), and cellular
[3 H]thymidine incorporation was measured by liquid
scintillation counting.

*Proliferative responses against syngeneic pancreatic
15 islets*

Islets of Langerhans from NOD mice were obtained as
previously described (Pankewycz, O., et al., 1991, Eur. J.
Immunol., 21:873), irradiated (3,000 rads) and placed in
flat-bottom 96-well plates with 10^5 T-cells in complete
20 RPMI. Cells were cultured for 6 days, pulsed with 1 μ Ci
[3 H]thymidine during the last 6 hours of culture, harvested
with a semiautomated cell harvester (PHD, Cambridge, MA),
and cellular incorporation of [3 H]thymidine was measured by
liquid scintillation counting.

25 Detection of suppressor activity

HT-2 or CTLL-2 cells were cultured at a
concentration of 10^5 cells/well in a final volume of 0.2 ml
complete RPMI media \pm rIL-2 \pm supernatants of NOD T-cell
clones and \pm anti-TGF- β 1 or anti-TGF- β 2 mAbs. After 20 hours
30 of culture, each well was pulsed with 1 μ Ci [3 H]thymidine,
incubated for an additional 4 hr, and harvested with a
semiautomated cell harvester (PHD, Cambridge, MA). Tritiated
thymidine incorporation was measured by liquid scintillation

counting. Suppressor activity was calculated according to the following formula: % suppressor activity: [experimental samples (supernatant + IL-2) - negative control (no IL-2)]/[positive control (IL-2 alone) - negative control].

- 5 Results were given as the mean of triplicate cultures. SEM did not exceed 12.5% of means. Similar results were found for HT-2 or CTLL-2 cells.

Bioassay for Transforming Growth Factor β (TGF- β)

- 10 Mink lung epithelial (CCL-64) TGF- β indicator cells were used as previously described (Danielpour, D., et al., 1989, J. Cell. Physiol., 138:79). Briefly, 10^5 cells/ml were placed in 24 well plates with culture media containing T-cell supernatants and cultured for 22 hours. The cells were then pulsed with 0.5 μ Ci/well of [3 H]thymidine for 2
15 hours, and [3 H]thymidine incorporation was measured by liquid scintillation counting. A TGF- β standard curve was constructed using TGF- β indicator cells cultured with serial dilutions of a known amount of TGF- β .

Flow cytometric analysis

- 20 HT-2 cells (2.5×10^5) were suspended in 50 μ l Hank's balanced salt solution containing 0.1% sodium azide and 20% (vol/vol) horse serum. The cells were incubated for 30 minutes at 4°C with 0.2 μ g of anti-IL-2R α mAb, washed and incubated with anti-rat IgG-FITC conjugated 30 minutes at
25 4°C. Subsequently, the cells were washed, resuspended in 2% paraformaldehyde and analyzed using a Becton and Dickinson FACS analyzer.

Example 11.

- Partial size purification. Supernatants harvested
30 from 24 hour cultures of IS-2.15 T-cells grown in complete RPMI without fetal bovine serum were loaded into 10, 30 and 100 KD membrane Centricon microconcentrator tubes (Amicon, Danvers, MA) with different membrane size cut-offs (10, 30

and 100 KD) and centrifuged at 6,000 RPM for 60 minutes at 25 °C. The concentrates and filtrates were tested for biological activity as described previously.

Example 12.

- 5 RNA extraction and Polymerase-Chain reaction (PCR)
procedure. Cytoplasmic RNA was extracted from T-cell clones
by the NP-40 lysis method. Two micrograms of RNA were
reverse transcribed into cDNA using random hexamer
oligo(dN)₆ as primer (BRL) and AMV reverse transcriptase
10 (Promega, Madison, WI) in a 50 µl reaction. Ten microliters
of the cDNA was amplified by PCR. PCR conditions implemented
in a 50 µl reaction were as follows: 75-750 pmol of each
primer (see below), 200 µM each dGTP, dATP, dCTP and dTTP
(Perkin-Elmer/Cetus, Emeryville, CA), 50 mM KCl, 10 mM Tris-
15 HCl, pH 8.3, 1.5 mM MgCl₂, and 2 units Taq DNA polymerase
("AmpliTaq", Perkin-Elmer/Cetus, Emeryville, CA). The
following primers were used: IL-2 sense primer
5'-TGATGGACCTACAGGAGCTCCTGAG-3' (nucleotides 203'-227') (SEQ
ID NO: 33); IL-2 antisense primer
20 5'-GAGTCAAATCCAGAAACATGCCGCAG-3' (nucleotides 370'-346')
(SEQ ID NO: 34); IL-4 sense primer
5'-CGAAGAACACCACAGAGAGTGAGCT-3' (nucleotides 231-255) (SEQ
ID NO: 35); IL-4 antisense primer 5'-
GACTCATTCATGGTGCAGCTTATCG-3' (nucleotides 411'-387') (SEQ ID
25 NO: 36); IL-6 sense primer 5'-TGGAGTCACAGAAGGAGTGGCTAAG-3'
(nucleotides 581'-605') (SEQ ID NO: 37); IL-6 antisense
primer 5'-TCTGACCACAGTGAGGAATGTCCAC-3' (nucleotides 735'-
711') (SEQ ID NO: 38); IFN-γ sense primer
5'-AGCGGCTGACTGAACTGAACTCAGATTGTAG-3' (nucleotides
30 841'-865') (SEQ ID NO: 39); IFN-γ antisense primer
5'-GTCACAGTTTTTCAGCTGTATAGGG-3' (nucleotides 1084'-1061')
(SEQ ID NO: 40); TNF-α sense primer
5'-GGCAGGTCTACTTTGGAGTCATTG-3' (nucleotides 820'-843') (SEQ

ID NO: 41); TNF- α antisense primer
5'-ACATTCGAGGCTCCAGTGAATTCCAG-3' (nucleotides 1127'-1102')
(SEQ ID NO: 42); TGF- β sense primer 5'-AAGTGGATCCACGAGCCCAA-
3' (nucleotides 1277'-1298') (SEQ ID NO: 43); TGF- β
5 antisense primer 5'-CTGCACTTGCAGGAGCGCAC-3' (nucleotides
1521'-1502') (SEQ ID NO: 44), as previously described
(Murray, L.J., et al., 1990, Eur. J. Immunol., 20:163).
Reactions were incubated in a Perkin-Elmer/Cetus DNA thermal
cycler for 45 cycles (denaturation 30 seconds, 95°C;
10 annealing 30 seconds, 56°C; extension 60 seconds, 72°C).
Fifteen microliters of the reaction product was
electrophoresed through a 1.2% agarose gel and transferred
onto a nylon membrane by capillary action (Zeta-probe, Bio-
Rad, Richmond, CA). Blots were hybridized at 65°C for 18
15 hours with 1 mM EDTA, 0.5 M NaH₂PO₄ (pH 7.2) and 7% sodium
dodecyl sulfate (SDS), and radiolabeled with probes for
murine rIL-2 (90 bp probe inserted in pBS) (Kasima, N., et
al., 1985, Nature, 313:401), IL-4 (1 kb probe inserted in
pXM) (Lee, F., et al., 1986, PNAS, 83:2063), IL-6 (1 kb
20 probe inserted in pXM) (Chiu, C.P., et al., 1988, PNAS,
85:7099), IFN- γ (643 bp probe inserted in pms10) (Gray,
P.W., et al., 1983, PNAS, 80:5842), TNF- α (480 bp probe
inserted in p-mTNF-1) (Fransen, L. et al., 1985, Nucl. Ac.
Res., 13:4417) and human TGF- β (2.14 kb probe inserted in
25 pBR327) (Derynck, R., et al., 1986, J. Biol. Chem.,
261:4377). After hybridization the blots were washed twice
at 65°C with 1 mM EDTA, 40 mM NaH₂PO₄ (pH 7.2) and 5% SDS
for 30 minutes, and twice with 1 mM EDTA, 40mM NaH₂PO₄ (pH
7.2) and 1% SDS for 30 minutes at 65°C. Blots were then
30 exposed to Kodak X-AR film at -70°C for 6-24 hours. Results
of the analysis of cytokine gene expression are described
below.

Example 13.

T-cell receptor (TCR) crosslinking did not induce proliferation in an IS-2.15 clone. To determine whether IS-2.15 is an anergic or normally responsive T-cell clone, culture plates were coated with mitogenic anti-TCR/CD3 complex and cellular proliferation was measured. Unlike NOD spleen cells, IS-2.15 T-cells did not proliferate in response to anti-CD3 or anti-V β 11 mAbs (Fig.5). Increasing the concentration of either the mAbs (Fig.5A and Fig.5B) or the concentration of responder cells (Fig.5C) failed to produce mitogenesis. These findings have been corroborated in each of three subsequent experiments. To determine whether this unresponsiveness to anti-TCR/CD3 mAbs was merely caused by a delay in proliferation the culture period was prolonged to 5 days. A failure to proliferate in response to anti-CD3 or anti-V β 11 mAbs was also noted at 5 days.

Example 14.

IS-2.15 clone proliferates, albeit slowly, to syngeneic spleen cells or pancreatic islets. To assess whether IS-2.15 T-cells are autoreactive 5×10^5 IS-2.15 T-cells/ml were co-cultured with 50-70 irradiated syngeneic NOD islets/ml or 10^6 spleen cells/ml in a final volume of 200 μ l. Although IS-2.15 T-cells responded with a far lower proliferative rate than many other organ-specific islet infiltrating T-cell clones (Pankewycz, O., et al., 1991, Eur. J. Immunol., 21:873), IS-2.15 T-cells do proliferate, albeit weakly, in response to islets. Table 3 shows a representative, i.e. one of three, experiment.

Example 16.

IS-2.15 supernatant specifically inhibits rIL-2 dependent Proliferation in rIL-2 dependent murine T-cell lines. To investigate the possibility that IS-2.15 T-cells

produce an immunosuppressive factor, the ability of IS-2.15 to block IL-2 stimulated proliferation of murine T-cells was tested. Two different murine IL-2 dependent T-cell lines, CTLL-2 and HT-2, were used as defined model systems for this analysis. When either of these T-cell lines are pre-incubated with IS-2.15 supernatant, their proliferative response to rIL-2 was abrogated, indicating that a suppressor or anti-proliferative substance is present in this supernatant (Fig.6A). To examine whether this inhibitory substance was produced by other islet infiltrating NOD T-cell clones, supernatants from two additional islet-infiltrating clones were analyzed. Neither clone produced a substance that inhibited the proliferative response of CTLL-2 or HT-2 cells to rIL-2. Supernatants from cloned G-3 cells did not modify rIL-2 stimulated proliferation, whereas supernatants from cloned H-4 cells actually amplified rIL-2 dependent T-cell proliferation. Titration of IS-2.15 suppressor activity showed that supernatant dilutions of 1/16 were suppressive (Fig.6B).

Although IL-2 indicator cells (CTLL-2 AND HT-2) cultured with IS-2.15 supernatant and rIL-2 do not proliferate, HT-2 cells remain viable after interaction with the supernatant as determined by dye trypan blue exclusion. Viability of cells cultured for 24 hours with rIL-2 was 99% with or without addition of IS-2.15 supernatant.

Example 16.

IS-2.15 supernatant inhibitory activity is not due to TGF- β . Since TGF- β is the only cytokine known to directly block the response of T-cells to IL-2, the capacity of neutralizing anti-TGF- β 1 and anti-TGF- β 2 antibodies to restore proliferation of HT-2 cells cultivated with rIL-2 and IS-2.15 T-cell supernatants was tested. Anti-TGF- β 1 and anti-TGF- β 2 antibodies, used at neutralizing concentrations

as described elsewhere (Danielpour, D., et al., 1989, J. Cell. Physiol., 138:79), failed to restore rIL-2 dependent proliferation of HT-2 cells. Moreover, TGF- β bioactivity was not detected in IS-2.15 T-cell supernatants using TGF- β dependent epithelial lines as indicator cells. It is doubtful that the active moiety is TGF- β .

Example 17.

Inhibition of rIL-2 stimulated proliferation of HT-2 cells is not readily reversible and is not due to downregulation of the IL-2 receptor (IL-2R α). Saturating concentrations of rIL-2 do not restore IL-2 dependent proliferation of HT-2 cells cultured with IS-2.15 supernatant (Fig.5A). The mechanism of this inhibitory activity was investigated. HT-2 cells were cultured with rIL-2 (50 u/ml) \pm IS-2.15 supernatant (10%, vol/vol) for 24 hours. HT-2 cells were washed twice and recultured for an additional 24 hours with different concentrations of rIL-2. HT-2 cells that were not pretreated with IS-2.15 supernatant responded to rIL-2. In contrast, HT-2 cells pre-cultured for 24 hours with IS-2.15 supernatant and washed did not proliferate in response to rIL-2. Since cell viability was not compromised by IS-2.15 supernatant (99% in the two groups), a cytotoxic effect is not responsible for the anti-proliferative activity (Fig. 7).

In theory, downregulation of IL-2R α expression could account for these effects. Therefore, IL-2R α expression was examined by flow cytometry after culturing HT-2 cells with rIL-2 (50 u/ml) \pm IS-2.15 supernatant (10%, vol/vol). IS-2.15 supernatant did not inhibit IL-2R α expression in HT-2 cells (Fig. 8).

Example 18.

IS-2.15 supernatant inhibitory factor is a heat-sensitive protein with a molecular weight between 10 and 30

KD. The inhibitory activity of IS-2.15 upon IL-2 dependent proliferation was completely destroyed by pre-heating IS-2.15 supernatant for 15 minutes at 65°C. Size selective ultracentrifugation procedures indicate that the molecular weight of this substance is between 10 and 30 KD (Fig. 9).

Example 19.

Analysis of cytokine gene expression by PCR.

Messenger RNA was extracted from each of the 3 islet-infiltrating NOD T-cell clones (IS-2.15, H-4, G-3) and amplified via PCR, as described above. TGF- β encoding mRNA was detected only in the IS-2.15 clone. This clone also expresses TNF- α mRNA. None of the clones expressed IL-2 encoding mRNA; IL-4 encoding mRNA was detected only in the H-4 islet-infiltrating T-cell clone. This finding correlated with the ability of the H-4 clone supernatant to increase proliferation of the IL-4 and IL-2 sensitive HT-2 line to rIL-2 (Fig. 6A). In contrast, the G-3 clone did not express IL-2 or IL-4 encoding mRNA. All 3 clones expressed IL-6 and IFN- γ mRNA.

Example 20.

Purification of the IL-2.15 suppressor factor

protein. The IS-2.15 suppressor factor protein or its component polypeptide(s) can be purified using conventional methods of protein purification known to one schooled in the art, e.g., methods including but not limited to precipitation, chromatography, immunoabsorption, or affinity techniques. The polypeptide can be purified from starting material using the IS-2.15 supernatant, an IS-2.15 suppressor factor cDNA or genomic DNA, as described below, or using a recombinant form of either of these DNAs genetically engineered into an overproducing cell line.

Example 21.

Isolation of a human IL-2.15 suppressor factor cDNA and genomic DNA. A cDNA encoding the IL-2.15 suppressor factor can be isolated from a human expression library by screening with antibodies to the suppressor factor.

- 5 Antibodies for screening can be raised in an animal, for example a rabbit. Possible immunogens include but are not limited to 1) a purified fragment of the suppressor factor protein obtained by conventional methods of protein separation as described above, or 2) a partially purified
- 10 sample of IL-2.15 suppressor factor. The protein sample can be partially purified by fractionating an IL-2.15 supernatant by size, as described above. Antibodies are labeled with a suitable label and are then used as probes to screen an expression library. The cDNA library can be
- 15 generated by isolating poly(A⁺) mRNA from the IL-2.15 cell line, converting the RNA to double stranded cDNA (Aruffo, A., et al., 1987, PNAS, 84:8573-7) and ligating the cDNA into a λ expression vector, for example λ gt11 (Clontech), or into the expression vector pCDNA-1 (Invitrogen, San Diego).
- 20 Positive clones can be confirmed by testing for the ability to block IL-2 stimulated T-cell proliferation.

The human suppressor factor gene can be cloned by analogous methods, for example by using a human cDNA library. The IL-2.15 cDNA sequences identified above can be

25 used as oligonucleotide probes to obtain the analogous human suppressor factor cDNA or genomic DNA by methods known to those skilled in the art.

SEQUENCE LISTING

(1) GENERAL INFORMATION:

- (i) APPLICANT: Strom, Terry
Rubin-Kelly, Vicki E.
Libermann, Towia
- (ii) TITLE OF INVENTION: METHODS AND COMPOUNDS FOR
PREVENTION OF GRAFT REJECTION
- (iii) NUMBER OF SEQUENCES: 32
- (iv) CORRESPONDENCE ADDRESS:
- (A) ADDRESSEE: Fish & Richardson
(B) STREET: 225 Franklin Street
(C) CITY: Boston
(D) STATE: Massachusetts
(E) COUNTRY: U.S.A.
(F) ZIP: 02110-2804
- (v) COMPUTER READABLE FORM:
- (A) MEDIUM TYPE: 3.5" Diskette, 1.44 Mb
(B) COMPUTER: IBM PS/2 Model 50Z or 55SX
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- (vi) CURRENT APPLICATION DATA:
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- (viii) ATTORNEY/AGENT INFORMATION:

(A) NAME: Paul T. Clark, Esq.
(B) REGISTRATION NUMBER: 30,162
(C) REFERENCE/DOCKET NUMBER: 05311/012001

(ix) TELECOMMUNICATION INFORMATION:

(A) TELEPHONE: (617) 542-5070
(B) TELEFAX: (617) 542-8906
(C) TELEX: 200154

(2) INFORMATION FOR SEQUENCE IDENTIFICATION NUMBER: 1:

(i) SEQUENCE CHARACTERISTICS:

(A) LENGTH: 25
(B) TYPE: nucleic acid
(C) STRANDEDNESS: single
(D) TOPOLOGY: linear

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 1:

TGATGGACCT ACAGGAGCTC CTGAG

25

(2) INFORMATION FOR SEQUENCE IDENTIFICATION NUMBER: 2:

(i) SEQUENCE CHARACTERISTICS:

(A) LENGTH: 26
(B) TYPE: nucleic acid
(C) STRANDEDNESS: single
(D) TOPOLOGY: linear

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 2:

GAGTCAAATC CAGAAACATG CCGCAG

26

(2) INFORMATION FOR SEQUENCE IDENTIFICATION NUMBER: 3:

(i) SEQUENCE CHARACTERISTICS:

(A) LENGTH: 25
(B) TYPE: nucleic acid
(C) STRANDEDNESS: single
(D) TOPOLOGY: linear

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 3:

CGAAGAACAC CACAGAGAGT GAGCT

25

(2) INFORMATION FOR SEQUENCE IDENTIFICATION NUMBER: 4:

(i) SEQUENCE CHARACTERISTICS:

(A) LENGTH:	25
(B) TYPE:	nucleic acid
(C) STRANDEDNESS:	single
(D) TOPOLOGY:	linear

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 4:

GACTCATTCA TGGTGCAGCT TATCG

25

(2) INFORMATION FOR SEQUENCE IDENTIFICATION NUMBER: 5:

(i) SEQUENCE CHARACTERISTICS:

(A) LENGTH:	25
(B) TYPE:	nucleic acid
(C) STRANDEDNESS:	single
(D) TOPOLOGY:	linear

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 5:

TGGAGTCACA GAAGGAGTGG CTAAG

25

(2) INFORMATION FOR SEQUENCE IDENTIFICATION NUMBER: 6:

(i) SEQUENCE CHARACTERISTICS:

(A) LENGTH:	25
(B) TYPE:	nucleic acid
(C) STRANDEDNESS:	single
(D) TOPOLOGY:	linear

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 6:

TCTGACCACA GTGAGGAATG TCCAC

25

(2) INFORMATION FOR SEQUENCE IDENTIFICATION NUMBER: 7:

(i) SEQUENCE CHARACTERISTICS:

(A) LENGTH: 31
(B) TYPE: nucleic acid
(C) STRANDEDNESS: single
(D) TOPOLOGY: linear

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 7:

AGCGGCTGAC TGAAGTGAAC TCAGATTGTA G 31

(2) INFORMATION FOR SEQUENCE IDENTIFICATION NUMBER: 8:

(i) SEQUENCE CHARACTERISTICS:

(A) LENGTH: 24
(B) TYPE: nucleic acid
(C) STRANDEDNESS: single
(D) TOPOLOGY: linear

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 8:

GTCACAGTTT TCAGCTGTAT AGGG 24

(2) INFORMATION FOR SEQUENCE IDENTIFICATION NUMBER: 9:

(i) SEQUENCE CHARACTERISTICS:

(A) LENGTH: 24
(B) TYPE: nucleic acid
(C) STRANDEDNESS: single
(D) TOPOLOGY: linear

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 9:

GGCAGGTCTA CTTTGGAGTC ATTG 24

(2) INFORMATION FOR SEQUENCE IDENTIFICATION NUMBER: 10:

(i) SEQUENCE CHARACTERISTICS:

(A) LENGTH: 26
(B) TYPE: nucleic acid
(C) STRANDEDNESS: single
(D) TOPOLOGY: linear

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 10:

ACATTCGAGG CTCCAGTGAA TTCCAG

26

(2) INFORMATION FOR SEQUENCE IDENTIFICATION NUMBER: 11:

(i) SEQUENCE CHARACTERISTICS:

(A) LENGTH: 20
(B) TYPE: nucleic acid
(C) STRANDEDNESS: single
(D) TOPOLOGY: linear

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 11:

AAGTGGATCC ACGAGCCCAA

20

(2) INFORMATION FOR SEQUENCE IDENTIFICATION NUMBER: 12:

(i) SEQUENCE CHARACTERISTICS:

(A) LENGTH: 20
(B) TYPE: nucleic acid
(C) STRANDEDNESS: single
(D) TOPOLOGY: linear

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 12:

CTGCACTTGC AGGAGCGCAC

20

(2) INFORMATION FOR SEQUENCE IDENTIFICATION NUMBER: 13:

(i) SEQUENCE CHARACTERISTICS:

(A) LENGTH: 30
(B) TYPE: nucleic acid
(C) STRANDEDNESS: single
(D) TOPOLOGY: linear

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 13:

CTTGGCATGC TTGTCAACAG CGCACCCACT

30

(2) INFORMATION FOR SEQUENCE IDENTIFICATION NUMBER: 14:

(i) SEQUENCE CHARACTERISTICS:

(A) LENGTH:	27
(B) TYPE:	nucleic acid
(C) STRANDEDNESS:	single
(D) TOPOLOGY:	linear

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 14:

GTGTTGTAAG CAGGAGGTAC ATAGTTA

27

(2) INFORMATION FOR SEQUENCE IDENTIFICATION NUMBER: 15:

(i) SEQUENCE CHARACTERISTICS:

(A) LENGTH:	22
(B) TYPE:	nucleic acid
(C) STRANDEDNESS:	single
(D) TOPOLOGY:	linear

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 15:

CACGGCACAG TCATTGAAAG CC

22

(2) INFORMATION FOR SEQUENCE IDENTIFICATION NUMBER: 16:

(i) SEQUENCE CHARACTERISTICS:

(A) LENGTH:	22
(B) TYPE:	nucleic acid
(C) STRANDEDNESS:	single
(D) TOPOLOGY:	linear

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 16:

TTCCGGCAAC AGCTGGTGGA CC

22

(2) INFORMATION FOR SEQUENCE IDENTIFICATION NUMBER: 17:

(i) SEQUENCE CHARACTERISTICS:

(A) LENGTH: 23
(B) TYPE: nucleic acid
(C) STRANDEDNESS: single
(D) TOPOLOGY: linear

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 17:

GATGTGCCAA ACGTCCTCAC AGC 23

(2) INFORMATION FOR SEQUENCE IDENTIFICATION NUMBER: 18:

(i) SEQUENCE CHARACTERISTICS:

(A) LENGTH: 24
(B) TYPE: nucleic acid
(C) STRANDEDNESS: single
(D) TOPOLOGY: linear

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 18:

CGATGAATCC AGGCATCGAA AAGC 24

(2) INFORMATION FOR SEQUENCE IDENTIFICATION NUMBER: 19:

(i) SEQUENCE CHARACTERISTICS:

(A) LENGTH: 22
(B) TYPE: nucleic acid
(C) STRANDEDNESS: single
(D) TOPOLOGY: linear

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 19:

CTGCCTGCTC TTACTGACTG GC 22

(2) INFORMATION FOR SEQUENCE IDENTIFICATION NUMBER: 20:

(i) SEQUENCE CHARACTERISTICS:

(A) LENGTH: 20
(B) TYPE: nucleic acid
(C) STRANDEDNESS: single
(D) TOPOLOGY: linear

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 20:

AATCACTCTT CACCTGCTCC

20

(2) INFORMATION FOR SEQUENCE IDENTIFICATION NUMBER: 21:

(i) SEQUENCE CHARACTERISTICS:

(A) LENGTH: 18
(B) TYPE: nucleic acid
(C) STRANDEDNESS: single
(D) TOPOLOGY: linear

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 21:

CCCAGGCGCA ATGTCAAT

18

(2) INFORMATION FOR SEQUENCE IDENTIFICATION NUMBER: 22:

(i) SEQUENCE CHARACTERISTICS:

(A) LENGTH: 18
(B) TYPE: nucleic acid
(C) STRANDEDNESS: single
(D) TOPOLOGY: linear

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 22:

CCAGGATAAG AAAC TCGA

18

(2) INFORMATION FOR SEQUENCE IDENTIFICATION NUMBER: 23:

(i) SEQUENCE CHARACTERISTICS:

(A) LENGTH: 20
(B) TYPE: nucleic acid

(C) STRANDEDNESS: single
(D) TOPOLOGY: linear

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 23:

AAGTGGATCC ACGAGCCCAA

20

(2) INFORMATION FOR SEQUENCE IDENTIFICATION NUMBER: 24:

(i) SEQUENCE CHARACTERISTICS:

(A) LENGTH: 20
(B) TYPE: nucleic acid
(C) STRANDEDNESS: single
(D) TOPOLOGY: linear

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 24:

CTGCACTTGC AGGAGCGCAC

20

(2) INFORMATION FOR SEQUENCE IDENTIFICATION NUMBER: 25:

(i) SEQUENCE CHARACTERISTICS:

(A) LENGTH: 21
(B) TYPE: nucleic acid
(C) STRANDEDNESS: single
(D) TOPOLOGY: linear

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 25:

GGTCTATATG CGTTGCTTAG G

21

(2) INFORMATION FOR SEQUENCE IDENTIFICATION NUMBER: 26:

(i) SEQUENCE CHARACTERISTICS:

(A) LENGTH: 20
(B) TYPE: nucleic acid
(C) STRANDEDNESS: single
(D) TOPOLOGY: linear

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 26:

CTCGGGAGAA GAATTTCTGC

20

(2) INFORMATION FOR SEQUENCE IDENTIFICATION NUMBER: 27:

(i) SEQUENCE CHARACTERISTICS:

(A) LENGTH:	24
(B) TYPE:	nucleic acid
(C) STRANDEDNESS:	single
(D) TOPOLOGY:	linear

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 27:

CGTGACATCA AAGAGAAGCT GTGC

24

(2) INFORMATION FOR SEQUENCE IDENTIFICATION NUMBER: 28:

(i) SEQUENCE CHARACTERISTICS:

(A) LENGTH:	25
(B) TYPE:	nucleic acid
(C) STRANDEDNESS:	single
(D) TOPOLOGY:	linear

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 28:

GCTCAGGAGG AGCAATGATC TTGAT

25

(2) INFORMATION FOR SEQUENCE IDENTIFICATION NUMBER: 29:

(i) SEQUENCE CHARACTERISTICS:

(A) LENGTH:	20
(B) TYPE:	nucleic acid
(C) STRANDEDNESS:	single
(D) TOPOLOGY:	linear

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 29:

ACCAGCCCTA AGTGATCCGC

20

(2) INFORMATION FOR SEQUENCE IDENTIFICATION NUMBER: 30:

(i) SEQUENCE CHARACTERISTICS:

(A) LENGTH: 24
(B) TYPE: nucleic acid
(C) STRANDEDNESS: single
(D) TOPOLOGY: linear

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 30:

GGTAGAGGGA GCAGATGCTG GTGC

24

(2) INFORMATION FOR SEQUENCE IDENTIFICATION NUMBER: 31:

(i) SEQUENCE CHARACTERISTICS:

(A) LENGTH: 24
(B) TYPE: nucleic acid
(C) STRANDEDNESS: single
(D) TOPOLOGY: linear

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 31:

GACCCTCAGG CCTACAAGGA GAGC

24

(2) INFORMATION FOR SEQUENCE IDENTIFICATION NUMBER: 32:

(i) SEQUENCE CHARACTERISTICS:

(A) LENGTH: 24
(B) TYPE: nucleic acid
(C) STRANDEDNESS: single
(D) TOPOLOGY: linear

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 32:

GGATCTCATA GAGGATGGTK GCAG

24

(2) INFORMATION FOR SEQUENCE IDENTIFICATION NUMBER: 33:

(i) SEQUENCE CHARACTERISTICS:

(A) LENGTH: 25
(B) TYPE: nucleic acid

(C) STRANDEDNESS: single
(D) TOPOLOGY: linear

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 33:

TGATGGACCT ACAGGAGCTC CTGAG

25

(2) INFORMATION FOR SEQUENCE IDENTIFICATION NUMBER: 34:

(i) SEQUENCE CHARACTERISTICS:

(A) LENGTH: 25
(B) TYPE: nucleic acid
(C) STRANDEDNESS: single
(D) TOPOLOGY: linear

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 34:

GAGTCAAATC CAGAAACATG CCGCAG

25

(2) INFORMATION FOR SEQUENCE IDENTIFICATION NUMBER: 35:

(i) SEQUENCE CHARACTERISTICS:

(A) LENGTH: 25
(B) TYPE: nucleic acid
(C) STRANDEDNESS: single
(D) TOPOLOGY: linear

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 35:

CGAAGAACAC CACAGAGAGT GAGCT

25

(2) INFORMATION FOR SEQUENCE IDENTIFICATION NUMBER: 36:

(i) SEQUENCE CHARACTERISTICS:

(A) LENGTH: 25
(B) TYPE: nucleic acid
(C) STRANDEDNESS: single
(D) TOPOLOGY: linear

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 36:

GACTCATTCA TGGTGCAGCT TATCG

25

(2) INFORMATION FOR SEQUENCE IDENTIFICATION NUMBER: 37:

(i) SEQUENCE CHARACTERISTICS:

(A) LENGTH: 25
(B) TYPE: nucleic acid
(C) STRANDEDNESS: single
(D) TOPOLOGY: linear

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 37:

TGGAGTCACA GAAGGAGTGG CTAAG

25

(2) INFORMATION FOR SEQUENCE IDENTIFICATION NUMBER: 38:

(i) SEQUENCE CHARACTERISTICS:

(A) LENGTH: 25
(B) TYPE: nucleic acid
(C) STRANDEDNESS: single
(D) TOPOLOGY: linear

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 38:

TCTGACCACA GTGAGGAATG TCCAC

25

(2) INFORMATION FOR SEQUENCE IDENTIFICATION NUMBER: 39:

(i) SEQUENCE CHARACTERISTICS:

(A) LENGTH: 31
(B) TYPE: nucleic acid
(C) STRANDEDNESS: single
(D) TOPOLOGY: linear

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 39:

AGCGGCTGAC TGAAGTGAAC TCAGATTGTA G

31

(2) INFORMATION FOR SEQUENCE IDENTIFICATION NUMBER: 40:

(i) SEQUENCE CHARACTERISTICS:

(A) LENGTH: 24
(B) TYPE: nucleic acid
(C) STRANDEDNESS: single
(D) TOPOLOGY: linear

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 40:

GTCACAGTTT TCAGCTGTAT AGGG

24

(2) INFORMATION FOR SEQUENCE IDENTIFICATION NUMBER: 41:

(i) SEQUENCE CHARACTERISTICS:

(A) LENGTH: 24
(B) TYPE: nucleic acid
(C) STRANDEDNESS: single
(D) TOPOLOGY: linear

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 41:

GGCAGGTCTA CTTTGGAGTC ATTG

24

(2) INFORMATION FOR SEQUENCE IDENTIFICATION NUMBER: 42:

(i) SEQUENCE CHARACTERISTICS:

(A) LENGTH: 26
(B) TYPE: nucleic acid
(C) STRANDEDNESS: single
(D) TOPOLOGY: linear

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 42:

ACATTCGAGG CTCCAGTGAA TTCCAG

26

(2) INFORMATION FOR SEQUENCE IDENTIFICATION NUMBER: 43:

(i) SEQUENCE CHARACTERISTICS:

(A) LENGTH: 21
(B) TYPE: nucleic acid

(C) STRANDEDNESS: single
(D) TOPOLOGY: linear

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 43:

AAGTGGATC CACGAGCCCA A

21

(2) INFORMATION FOR SEQUENCE IDENTIFICATION NUMBER: 44:

(i) SEQUENCE CHARACTERISTICS:

(A) LENGTH: 20
(B) TYPE: nucleic acid
(C) STRANDEDNESS: single
(D) TOPOLOGY: linear

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 44:

CTGCACTTGC AGGAGCGCAC

20